



Sequence Listing

<110> Innogenetics N.V.

<120> Purified hepatitis C virus envelope proteins for diagnostic and therapeutic use.

<130> 2551-69

<140> 09/995,860

<160> 122

<170> PatentIn 3.1

<210> 1

<211> 21

<212> DNA

<213> Hepatitis C virus

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21

<210> 2

<211> 68

<212> DNA

<213> Hepatitis C virus

<400> 2

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<213> Hepatitis C virus

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<221> CDS

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<221> mat_peptide

<222> 1..636

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48

ctg acc att cca gct tcc gct tat gag gtg cgc aac gtg tcc ggg atg
Leu Thr Ile Pro Ala Ser Ala Tyr Glu Val Arg Asn Val Ser Gly Met
20 25 30

96

tac cat gtc acg aac gac tgc tcc aac tca agc att gtg tat gag gca 144
 Tyr His Val Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr Glu Ala
 35 40 45

gcg gac atg atc atg cac acc ccc ggg tgc gtg ccc tgc gtt cgg gag 192
 Ala Asp Met Ile Met His Thr Pro Gly Cys Val Pro Cys Val Arg Glu
 50 55 60

aac aac tct tcc cgc tgc tgg gta gcg ctc acc ccc acg ctc gca gct 240
 Asn Asn Ser Ser Arg Cys Trp Val Ala Leu Thr Pro Thr Leu Ala Ala
 65 70 75 80

agg aac gcc agc gtc ccc acc acg aca ata cga cgc cac gtc gat ttg 288
 Arg Asn Ala Ser Val Pro Thr Thr Thr Ile Arg Arg His Val Asp Leu
 85 90 95

ctc gtt ggg gcg gct gct ctc tgt tcc gct atg tac gtg ggg gat ctc 336
 Leu Val Gly Ala Ala Leu Cys Ser Ala Met Tyr Val Gly Asp Leu
 100 105 110

tgc gga tct gtc ttc ctc gtc tcc cag ctg ttc acc atc tcg cct cgc 384
 Cys Gly Ser Val Phe Leu Val Ser Gln Leu Phe Thr Ile Ser Pro Arg
 115 120 125

cgg cat gag acg gtg cag gac tgc aat tgc tca atc tat ccc ggc cac 432
 Arg His Glu Thr Val Gln Asp Cys Asn Cys Ser Ile Tyr Pro Gly His
 130 135 140

ata aca ggt cac cgt atg gct tgg gat atg atg atg aac tgg tcg cct 480
 Ile Thr Gly His Arg Met Ala Trp Asp Met Met Met Asn Trp Ser Pro
 145 150 155 160

aca acg gcc ctg gtg gta tcg cag ctg ctc cgg atc cca caa gct gtc 528
 Thr Thr Ala Leu Val Val Ser Gln Leu Leu Arg Ile Pro Gln Ala Val
 165 170 175

gtg gac atg gtg gcg ggg gcc cat tgg gga gtc ctg gcg ggc ctc gcc 576
 Val Asp Met Val Ala Gly Ala His Trp Gly Val Leu Ala Gly Leu Ala
 180 185 190

tac tat tcc atg gtg ggg aac tgg gct aag gtt ttg att gtg atg cta 624
 Tyr Tyr Ser Met Val Gly Asn Trp Ala Lys Val Leu Ile Val Met Leu
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<211> 212

<212> PRT

<213> Hepatitis C virus

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 35 40 45
 Ala Asp Met Ile Met His Thr Pro Gly Cys Val Pro Cys Val Arg Glu
 50 55 60
 Asn Asn Ser Ser Arg Cys Trp Val Ala Leu Thr Pro Thr Leu Ala Ala
 65 70 75 80
 Arg Asn Ala Ser Val Pro Thr Thr Thr Ile Arg Arg His Val Asp Leu
 85 90 95
 Leu Val Gly Ala Ala Ala Leu Cys Ser Ala Met Tyr Val Gly Asp Leu
 100 105 110
 Cys Gly Ser Val Phe Leu Val Ser Gln Leu Phe Thr Ile Ser Pro Arg
 115 120 125
 Arg His Glu Thr Val Gln Asp Cys Asn Cys Ser Ile Tyr Pro Gly His
 130 135 140
 Ile Thr Gly His Arg Met Ala Trp Asp Met Met Met Asn Trp Ser Pro
 145 150 155 160
 Thr Thr Ala Leu Val Val Ser Gln Leu Leu Arg Ile Pro Gln Ala Val
 165 170 175
 Val Asp Met Val Ala Gly Ala His Trp Gly Val Leu Ala Gly Leu Ala
 180 185 190
 Tyr Tyr Ser Met Val Gly Asn Trp Ala Lys Val Leu Ile Val Met Leu
 195 200 205
 Leu Phe Ala Leu
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<212> DNA

<213> Hepatitis C virus

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 gtg ggg tac att ccg ctc gtc ggc gcc ccc cta ggg ggc gct gcc agg 96
 Val Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu Gly Gly Ala Ala Arg
 20 25 30

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gcc ctg gcg cat ggc gtc cgg gtt ctg gag gac ggc gtg aac tat gca      144
Ala Leu Ala His Gly Val Arg Val Leu Glu Asp Gly Val Asn Tyr Ala
      35              40              45

aca ggg aat ttg ccc ggt tgc tct ttc tct atc ttc ctc ttg gct ttg      192
Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu
      50              55              60

ctg tcc tgt ctg acc gtt cca gct tcc gct tat gaa gtg cgc aac gtg      240
Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Tyr Glu Val Arg Asn Val
      65              70              75              80

tcc ggg atg tac cat gtc acg aac gac tgc tcc aac tca agc att gtg      288
Ser Gly Met Tyr His Val Thr Asn Asp Cys Ser Asn Ser Ser Ile Val
      85              90              95

tat gag gca gcg gac atg atc atg cac acc ccc ggg tgc gtg ccc tgc      336
Tyr Glu Ala Ala Asp Met Ile Met His Thr Pro Gly Cys Val Pro Cys
      100             105             110

gtt cgg gag aac aac tct tcc cgc tgc tgg gta gcg ctc acc ccc acg      384
Val Arg Glu Asn Asn Ser Ser Arg Cys Trp Val Ala Leu Thr Pro Thr
      115             120             125

ctc gca gct agg aac gcc agc gtc ccc acc acg aca ata cga cgc cac      432
Leu Ala Ala Arg Asn Ala Ser Val Pro Thr Thr Thr Ile Arg Arg His
      130             135             140

gtc gat ttg ctc gtt ggg gcg gct gct ttc tgt tcc gct atg tac gtg      480
Val Asp Leu Leu Val Gly Ala Ala Phe Cys Ser Ala Met Tyr Val
      145             150             155             160

ggg gac ctc tgc gga tct gtc ttc ctc gtc tcc cag ctg ttc acc atc      528
Gly Asp Leu Cys Gly Ser Val Phe Leu Val Ser Gln Leu Phe Thr Ile
      165             170             175

tcg cct cgc cgg cat gag acg gtg cag gac tgc aat tgc tca atc tat      576
Ser Pro Arg Arg His Glu Thr Val Gln Asp Cys Asn Cys Ser Ile Tyr
      180             185             190

ccc ggc cac ata acg ggt cac cgt atg gct tgg gat atg atg atg aac      624
Pro Gly His Ile Thr Gly His Arg Met Ala Trp Asp Met Met Met Asn
      195             200             205

tgg tcg cct aca acg gcc ctg gtg gta tcg cag ctg ctc cgg atc cca      672
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      210             215             220

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Gln Ala Val Val Asp Met Val Ala Gly Ala His Trp Gly Val Leu Ala
      225             230             235             240

ggg ctc gcc tac tat tcc atg gtg ggg aac tgg gct aag gtt ttg att      768
Gly Leu Ala Tyr Tyr Ser Met Val Gly Asn Trp Ala Lys Val Leu Ile
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<212> PRT

<213> Hepatitis C virus

<400> 6

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Ala Leu Ala His Gly Val Arg Val Leu Glu Asp Gly Val Asn Tyr Ala
          35           40           45

Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu
          50           55           60

Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Tyr Glu Val Arg Asn Val
          65           70           75           80

Ser Gly Met Tyr His Val Thr Asn Asp Cys Ser Asn Ser Ser Ile Val
          85           90           95

Tyr Glu Ala Ala Asp Met Ile Met His Thr Pro Gly Cys Val Pro Cys
          100          105          110

Val Arg Glu Asn Asn Ser Ser Arg Cys Trp Val Ala Leu Thr Pro Thr
          115          120          125

Leu Ala Ala Arg Asn Ala Ser Val Pro Thr Thr Thr Ile Arg Arg His
          130          135          140

Val Asp Leu Leu Val Gly Ala Ala Ala Phe Cys Ser Ala Met Tyr Val
          145          150          155          160

Gly Asp Leu Cys Gly Ser Val Phe Leu Val Ser Gln Leu Phe Thr Ile
          165          170          175

Ser Pro Arg Arg His Glu Thr Val Gln Asp Cys Asn Cys Ser Ile Tyr
          180          185          190

Pro Gly His Ile Thr Gly His Arg Met Ala Trp Asp Met Met Met Asn
          195          200          205

Trp Ser Pro Thr Thr Ala Leu Val Val Ser Gln Leu Leu Arg Ile Pro
          210          215          220

Gln Ala Val Val Asp Met Val Ala Gly Ala His Trp Gly Val Leu Ala
          225          230          235          240

Gly Leu Ala Tyr Tyr Ser Met Val Gly Asn Trp Ala Lys Val Leu Ile
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Val Met Leu Leu Phe Ala Pro
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Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu Gly Gly Ala Ala Arg	
20 25 30	
gcc ctg gcg cat ggc gtc cgg gtt ctg gaa gac ggc gtg aac tat gca	144
Ala Leu Ala His Gly Val Arg Val Leu Glu Asp Gly Val Asn Tyr Ala	
35 40 45	
aca ggg aat ttg cct ggt tgc tct ttc tct atc ttc ctc ttg gct tta	192
Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu	
50 55 60	
ctg tcc tgt ctg acc att cca gct tcc gct tat gag gtg cgc aac gtg	240
Leu Ser Cys Leu Thr Ile Pro Ala Ser Ala Tyr Glu Val Arg Asn Val	
65 70 75 80	
tcc ggg atg tac cat gtc acg aac gac tgc tcc aac tca agc att gtg	288
Ser Gly Met Tyr His Val Thr Asn Asp Cys Ser Asn Ser Ser Ile Val	
85 90 95	
tat gag gca gcg gac atg atc atg cac acc ccc ggg tgc gtg ccc tgc	336
Tyr Glu Ala Ala Asp Met Ile Met His Thr Pro Gly Cys Val Pro Cys	
100 105 110	
gtt cgg gag aac aac tct tcc cgc tgc tgg gta gcg ctc acc ccc acg	384
Val Arg Glu Asn Asn Ser Ser Arg Cys Trp Val Ala Leu Thr Pro Thr	
115 120 125	
ctc gca gct agg aac gcc agc gtc ccc act acg aca ata cga cgc cac	432
Leu Ala Ala Arg Asn Ala Ser Val Pro Thr Thr Thr Ile Arg Arg His	
130 135 140	
gtc gat ttg ctc gtt ggg gcg gct gct ttc tgt tcc gct atg tac gtg	480
Val Asp Leu Leu Val Gly Ala Ala Ala Phe Cys Ser Ala Met Tyr Val	
145 150 155 160	
ggg gat ctc tgc gga tct gtc ttc ctc gtc tcc cag ctg ttc acc atc	528
Gly Asp Leu Cys Gly Ser Val Phe Leu Val Ser Gln Leu Phe Thr Ile	
165 170 175	
tcg cct cgc cgg cat gag acg gtg cag gac tgc aat tgc tca atc tat	576
Ser Pro Arg Arg His Glu Thr Val Gln Asp Cys Asn Cys Ser Ile Tyr	
180 185 190	
ccc ggc cac ata aca ggt cac cgt atg gct tgg gat atg atg atg aac	624
Pro Gly His Ile Thr Gly His Arg Met Ala Trp Asp Met Met Met Asn	
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tgg taatag	633
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<213> Hepatitis C virus

<400> 8

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Ala Leu Ala His Gly Val Arg Val Leu Glu Asp Gly Val Asn Tyr Ala
      35             40             45

Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu
      50             55             60

Leu Ser Cys Leu Thr Ile Pro Ala Ser Ala Tyr Glu Val Arg Asn Val
      65             70             75             80

Ser Gly Met Tyr His Val Thr Asn Asp Cys Ser Asn Ser Ser Ile Val
      85             90             95

Tyr Glu Ala Ala Asp Met Ile Met His Thr Pro Gly Cys Val Pro Cys
      100            105            110

Val Arg Glu Asn Asn Ser Ser Arg Cys Trp Val Ala Leu Thr Pro Thr
      115            120            125

Leu Ala Ala Arg Asn Ala Ser Val Pro Thr Thr Thr Ile Arg Arg His
      130            135            140

Val Asp Leu Leu Val Gly Ala Ala Ala Phe Cys Ser Ala Met Tyr Val
      145            150            155            160

Gly Asp Leu Cys Gly Ser Val Phe Leu Val Ser Gln Leu Phe Thr Ile
      165            170            175

Ser Pro Arg Arg His Glu Thr Val Gln Asp Cys Asn Cys Ser Ile Tyr
      180            185            190

Pro Gly His Ile Thr Gly His Arg Met Ala Trp Asp Met Met Met Asn
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ctg acc ata cca gct tcc gct tat gaa gtg cgc aac gtg tcc ggg gtg	96
Leu Thr Ile Pro Ala Ser Ala Tyr Glu Val Arg Asn Val Ser Gly Val	
20 25 30	
tac cat gtc acg aac gac tgc tcc aac tca agc ata gtg tat gag gca	144
Tyr His Val Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr Glu Ala	
35 40 45	
gcg gac atg atc atg cac acc ccc ggg tgc gtg ccc tgc gtt cgg gag	192
Ala Asp Met Ile Met His Thr Pro Gly Cys Val Pro Cys Val Arg Glu	
50 55 60	
ggc aac tcc tcc cgt tgc tgg gtg gcg ctc act ccc acg ctc gcg gcc	240
Gly Asn Ser Ser Arg Cys Trp Val Ala Leu Thr Pro Thr Leu Ala Ala	
65 70 75 80	
agg aac gcc agc gtc ccc aca acg aca ata cga cgc cac gtc gat ttg	288
Arg Asn Ala Ser Val Pro Thr Thr Thr Ile Arg Arg His Val Asp Leu	
85 90 95	
ctc gtt ggg gct gct gct ttc tgt tcc gct atg tac gtg ggg gat ctc	336
Leu Val Gly Ala Ala Ala Phe Cys Ser Ala Met Tyr Val Gly Asp Leu	
100 105 110	
tgc gga tct gtt ttc ctt gtt tcc cag ctg ttc acc ttc tca cct cgc	384
Cys Gly Ser Val Phe Leu Val Ser Gln Leu Phe Thr Phe Ser Pro Arg	
115 120 125	
cgg cat caa aca gta cag gac tgc aac tgc tca atc tat ccc ggc cat	432
Arg His Gln Thr Val Gln Asp Cys Asn Cys Ser Ile Tyr Pro Gly His	
130 135 140	
gta tca ggt cac cgc atg gct tgg gat atg atg atg aac tgg tcc taatag	483
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Arg Asn Ala Ser Val Pro Thr Thr Thr Ile Arg Arg His Val Asp Leu
85 90 95

ctc gtt ggg gct gct gct ttc tgt tcc gct atg tac gtg ggg gat ctc 336
 Leu Val Gly Ala Ala Ala Phe Cys Ser Ala Met Tyr Val Gly Asp Leu
 100 105 110

tgc gga tct gtt ttc ctt gtt tcc cag ctg ttc acc ttc tca cct cgc 384
 Cys Gly Ser Val Phe Leu Val Ser Gln Leu Phe Thr Phe Ser Pro Arg
 115 120 125

cgg cat caa aca gta cag gac tgc aac tgc tca atc tat ccc ggc cat 432
 Arg His Gln Thr Val Gln Asp Cys Asn Cys Ser Ile Tyr Pro Gly His
 130 135 140

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 35 40 45

Ala Asp Met Ile Met His Thr Pro Gly Cys Val Pro Cys Val Arg Glu
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Gly Asn Ser Ser Arg Cys Trp Val Ala Leu Thr Pro Thr Leu Ala Ala
 65 70 75 80

Arg Asn Ala Ser Val Pro Thr Thr Thr Ile Arg Arg His Val Asp Leu
 85 90 95

Leu Val Gly Ala Ala Ala Phe Cys Ser Ala Met Tyr Val Gly Asp Leu
 100 105 110

Cys Gly Ser Val Phe Leu Val Ser Gln Leu Phe Thr Phe Ser Pro Arg
 115 120 125

Arg His Gln Thr Val Gln Asp Cys Asn Cys Ser Ile Tyr Pro Gly His
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Val Ser Gly His Arg Met Ala Trp Asp Met Met Met Asn Trp
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<210> 13

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gtg ggg tac att ccg ctc gtc ggc gcc ccc cta ggg ggc gct gcc agg      96
Val Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu Gly Gly Ala Ala Arg
          20          25          30

gcc ctg gcg cat ggc gtc cgg gtt ctg gaa gac ggc gtg aac tat gca      144
Ala Leu Ala His Gly Val Arg Val Leu Glu Asp Gly Val Asn Tyr Ala
          35          40          45

aca ggg aat ttg cct ggt tgc tct ttc tct atc ttc ctc ttg gct tta      192
Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu
          50          55          60
ctg tcc tgt cta acc att cca gct tcc gct tac gag gtg cgc aac gtg      240
Leu Ser Cys Leu Thr Ile Pro Ala Ser Ala Tyr Glu Val Arg Asn Val
          65          70          75          80

tcc ggg atg tac cat gtc acg aac gac tgc tcc aac tca agc att gtg      288
Ser Gly Met Tyr His Val Thr Asn Asp Cys Ser Asn Ser Ser Ile Val
          85          90          95

tat gag gca gcg gac atg atc atg cac acc ccc ggg tgc gtg ccc tgc      336
Tyr Glu Ala Ala Asp Met Ile Met His Thr Pro Gly Cys Val Pro Cys
          100          105          110

gtt cgg gag aac aac tct tcc cgc tgc tgg gta gcg ctc acc ccc acg      384
Val Arg Glu Asn Asn Ser Ser Arg Cys Trp Val Ala Leu Thr Pro Thr
          115          120          125

ctc gcg gct agg aac gcc agc atc ccc act aca aca ata cga cgc cac      432
Leu Ala Ala Arg Asn Ala Ser Ile Pro Thr Thr Thr Ile Arg Arg His
          130          135          140

gtc gat ttg ctc gtt ggg gcg gct gct ttc tgt tcc gct atg tac gtg      480
Val Asp Leu Leu Val Gly Ala Ala Ala Phe Cys Ser Ala Met Tyr Val
          145          150          155          160

ggg gat ctc tgc gga tct gtc ttc ctc gtc tcc cag ctg ttc acc atc      528
Gly Asp Leu Cys Gly Ser Val Phe Leu Val Ser Gln Leu Phe Thr Ile
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tcg cct cgc cgg cat gag acg gtg cag gac tgc aat tgc tca atc tat      576
Ser Pro Arg Arg His Glu Thr Val Gln Asp Cys Asn Cys Ser Ile Tyr
          180          185          190

ccc ggc cac ata acg ggt cac cgt atg gct tgg gat atg atg atg aac      624
Pro Gly His Ile Thr Gly His Arg Met Ala Trp Asp Met Met Met Asn
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tgg tac taatag      636
Trp Tyr
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Ala Leu Ala His Gly Val Arg Val Leu Glu Asp Gly Val Asn Tyr Ala
      35             40             45

Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu
      50             55             60

Leu Ser Cys Leu Thr Ile Pro Ala Ser Ala Tyr Glu Val Arg Asn Val
      65             70             75             80

Ser Gly Met Tyr His Val Thr Asn Asp Cys Ser Asn Ser Ser Ile Val
      85             90             95

Tyr Glu Ala Ala Asp Met Ile Met His Thr Pro Gly Cys Val Pro Cys
      100            105            110

Val Arg Glu Asn Asn Ser Ser Arg Cys Trp Val Ala Leu Thr Pro Thr
      115            120            125

Leu Ala Ala Arg Asn Ala Ser Ile Pro Thr Thr Thr Ile Arg Arg His
      130            135            140

Val Asp Leu Leu Val Gly Ala Ala Ala Phe Cys Ser Ala Met Tyr Val
      145            150            155            160

Gly Asp Leu Cys Gly Ser Val Phe Leu Val Ser Gln Leu Phe Thr Ile
      165            170            175

Ser Pro Arg Arg His Glu Thr Val Gln Asp Cys Asn Cys Ser Ile Tyr
      180            185            190

Pro Gly His Ile Thr Gly His Arg Met Ala Trp Asp Met Met Met Asn
      195            200            205

Trp Tyr
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<210> 18

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<212> DNA

<213> Hepatitis C virus

<400> 18

ctattaccag ttcacatca tatccca

27

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<212> DNA

<213> Hepatitis C virus

<400> 19

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<221> misc_feature

<223>/note="antisense"

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gtg ggg tac att ccg ctc gtc ggc gcc ccc cta ggg ggc gct gcc agg	96
Val Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu Gly Gly Ala Ala Arg	
20 25 30	
gcc ctg gcg cat ggc gtc cgg gtt ctg gag gac ggc gtg aac tat gca	144
Ala Leu Ala His Gly Val Arg Val Leu Glu Asp Gly Val Asn Tyr Ala	
35 40 45	
aca ggg aat ttg ccc ggt tgc tct ttc tct atc ttc ctc ttg gct ttg	192
Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu	
50 55 60	
ctg tcc tgt ctg acc gtt cca gct tcc gct tat gaa gtg cgc aac gtg	240
Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Tyr Glu Val Arg Asn Val	
65 70 75 80	
tcc ggg atg tac cat gtc acg aac gac tgc tcc aac tca agc att gtg	288
Ser Gly Met Tyr His Val Thr Asn Asp Cys Ser Asn Ser Ser Ile Val	
85 90 95	
tat gag gca gcg gac atg atc atg cac acc ccc ggg tgc gtg ccc tgc	336
Tyr Glu Ala Ala Asp Met Ile Met His Thr Pro Gly Cys Val Pro Cys	
100 105 110	
gtt cgg gag aac aac tct tcc cgc tgc tgg gta gcg ctc acc ccc acg	384
Val Arg Glu Asn Asn Ser Ser Arg Cys Trp Val Ala Leu Thr Pro Thr	
115 120 125	
ctc gca gct agg aac gcc agc gtc ccc acc acg aca ata cga cgc cac	432
Leu Ala Ala Arg Asn Ala Ser Val Pro Thr Thr Thr Ile Arg Arg His	
130 135 140	
gtc gat tcc cag ctg ttc acc atc tcg cct cgc cgg cat gag acg gtg	480
Val Asp Ser Gln Leu Phe Thr Ile Ser Pro Arg Arg His Glu Thr Val	
145 150 155 160	
cag gac tgc aat tgc tca atc tat ccc ggc cac ata acg ggt cac cgt	528
Gln Asp Cys Asn Cys Ser Ile Tyr Pro Gly His Ile Thr Gly His Arg	
165 170 175	
atg gct tgg gat atg atg atg aac tgg tcg cct aca acg gcc ctg gtg	576
Met Ala Trp Asp Met Met Met Asn Trp Ser Pro Thr Thr Ala Leu Val	
180 185 190	
gta tcg cag ctg ctc cgg atc cca caa gct gtc gtg gac atg gtg gcg	624
Val Ser Gln Leu Leu Arg Ile Pro Gln Ala Val Val Asp Met Val Ala	
195 200 205	

ggg gcc cat tgg gga gtc ctg gcg ggt ctc gcc tac tat tcc atg gtg 672
 Gly Ala His Trp Gly Val Leu Ala Gly Leu Ala Tyr Tyr Ser Met Val
 210 215 220

ggg aac tgg gct aag gtt ttg att gtg atg cta ctc ttt gct ccc taatag 723
 Gly Asn Trp Ala Lys Val Leu Ile Val Met Leu Leu Phe Ala Pro
 225 230 235

<210> 22

<211> 239

<212> PRT

<213> Hepatitis C virus

<400> 22

Met Leu Gly Lys Val Ile Asp Thr Leu Thr Cys Gly Phe Ala Asp Leu
 1 5 10 15

Val Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu Gly Gly Ala Ala Arg
 20 25 30

Ala Leu Ala His Gly Val Arg Val Leu Glu Asp Gly Val Asn Tyr Ala
 35 40 45

Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu
 50 55 60

Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Tyr Glu Val Arg Asn Val
 65 70 75 80

Ser Gly Met Tyr His Val Thr Asn Asp Cys Ser Asn Ser Ser Ile Val
 85 90 95

Tyr Glu Ala Ala Asp Met Ile Met His Thr Pro Gly Cys Val Pro Cys
 100 105 110

Val Arg Glu Asn Asn Ser Ser Arg Cys Trp Val Ala Leu Thr Pro Thr
 115 120 125

Leu Ala Ala Arg Asn Ala Ser Val Pro Thr Thr Thr Ile Arg Arg His
 130 135 140

Val Asp Ser Gln Leu Phe Thr Ile Ser Pro Arg Arg His Glu Thr Val
 145 150 155 160

Gln Asp Cys Asn Cys Ser Ile Tyr Pro Gly His Ile Thr Gly His Arg
 165 170 175

Met Ala Trp Asp Met Met Met Asn Trp Ser Pro Thr Thr Ala Leu Val
 180 185 190

Val Ser Gln Leu Leu Arg Ile Pro Gln Ala Val Val Asp Met Val Ala
 195 200 205

Gly Ala His Trp Gly Val Leu Ala Gly Leu Ala Tyr Tyr Ser Met Val
 210 215 220

Gly Asn Trp Ala Lys Val Leu Ile Val Met Leu Leu Phe Ala Pro
 225 230 235

<210> 23

<211> 561
 <212> DNA
 <213> Hepatitis C virus

<220>
 <221> CDS
 <222> 1..558

<220>
 <221> mat_peptide
 <222> 1..555

<400> 23

atg	ttg	ggt	aag	gtc	atc	gat	acc	ctt	aca	tgc	ggc	ttc	gcc	gac	ctc	48
Met	Leu	Gly	Lys	Val	Ile	Asp	Thr	Leu	Thr	Cys	Gly	Phe	Ala	Asp	Leu	
1				5				10					15			

gtg	ggg	tac	att	ccg	ctc	gtc	ggc	gcc	ccc	cta	ggg	ggc	gct	gcc	agg	96
Val	Gly	Tyr	Ile	Pro	Leu	Val	Gly	Ala	Pro	Leu	Gly	Gly	Ala	Ala	Arg	
			20					25					30			

gcc	ctg	gcg	cat	ggc	gtc	cgg	gtt	ctg	gag	gac	ggc	gtg	aac	tat	gca	144
Ala	Leu	Ala	His	Gly	Val	Arg	Val	Leu	Glu	Asp	Gly	Val	Asn	Tyr	Ala	
		35					40					45				

aca	ggg	aat	ttg	ccc	ggg	tgc	tct	ttc	tct	atc	ttc	ctc	ttg	gct	ttg	192
Thr	Gly	Asn	Leu	Pro	Gly	Cys	Ser	Phe	Ser	Ile	Phe	Leu	Leu	Ala	Leu	
	50					55					60					

ctg	tcc	tgt	ctg	acc	gtt	cca	gct	tcc	gct	tat	gaa	gtg	cgc	aac	gtg	240
Leu	Ser	Cys	Leu	Thr	Val	Pro	Ala	Ser	Ala	Tyr	Glu	Val	Arg	Asn	Val	
65					70					75					80	

tcc	ggg	atg	tac	cat	gtc	acg	aac	gac	tgc	tcc	aac	tca	agc	att	gtg	288
Ser	Gly	Met	Tyr	His	Val	Thr	Asn	Asp	Cys	Ser	Asn	Ser	Ser	Ile	Val	
			85					90						95		

tat	gag	gca	gcg	gac	atg	atc	atg	cac	acc	ccc	ggg	tgc	gtg	ccc	tgc	336
Tyr	Glu	Ala	Ala	Asp	Met	Ile	Met	His	Thr	Pro	Gly	Cys	Val	Pro	Cys	
			100					105					110			

gtt	cgg	gag	aac	aac	tct	tcc	cgc	tgc	tgg	gta	gcg	ctc	acc	ccc	acg	384
Val	Arg	Glu	Asn	Asn	Ser	Ser	Arg	Cys	Trp	Val	Ala	Leu	Thr	Pro	Thr	
	115						120					125				

ctc	gca	gct	agg	aac	gcc	agc	gtc	ccc	acc	acg	aca	ata	cga	cgc	cac	432
Leu	Ala	Ala	Arg	Asn	Ala	Ser	Val	Pro	Thr	Thr	Thr	Ile	Arg	Arg	His	
	130						135					140				

gtc	gat	tcc	cag	ctg	ttc	acc	atc	tgc	cct	cgc	cgg	cat	gag	acg	gtg	480
Val	Asp	Ser	Gln	Leu	Phe	Thr	Ile	Ser	Pro	Arg	Arg	His	Glu	Thr	Val	
145					150				155					160		

cag	gac	tgc	aat	tgc	tca	atc	tat	ccc	ggc	cac	ata	acg	ggg	cac	cgt	528
Gln	Asp	Cys	Asn	Cys	Ser	Ile	Tyr	Pro	Gly	His	Ile	Thr	Gly	His	Arg	
			165					170						175		

atg	gct	tgg	gat	atg	atg	atg	aac	tgg	taatag							561
Met	Ala	Trp	Asp	Met	Met	Met	Asn	Trp								
			180					185								

<210> 24

<211> 185
 <212> PRT
 <213> Hepatitis C virus

<400> 24

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Met Leu Gly Lys Val Ile Asp Thr Leu Thr Cys Gly Phe Ala Asp Leu
 1             5             10             15

Val Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu Gly Gly Ala Ala Arg
      20             25             30

Ala Leu Ala His Gly Val Arg Val Leu Glu Asp Gly Val Asn Tyr Ala
      35             40             45

Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu
      50             55             60

Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Tyr Glu Val Arg Asn Val
      65             70             75             80

Ser Gly Met Tyr His Val Thr Asn Asp Cys Ser Asn Ser Ser Ile Val
      85             90             95

Tyr Glu Ala Ala Asp Met Ile Met His Thr Pro Gly Cys Val Pro Cys
      100            105            110

Val Arg Glu Asn Asn Ser Ser Arg Cys Trp Val Ala Leu Thr Pro Thr
      115            120            125

Leu Ala Ala Arg Asn Ala Ser Val Pro Thr Thr Thr Ile Arg Arg His
      130            135            140

Val Asp Ser Gln Leu Phe Thr Ile Ser Pro Arg Arg His Glu Thr Val
      145            150            155            160

Gln Asp Cys Asn Cys Ser Ile Tyr Pro Gly His Ile Thr Gly His Arg
      165            170            175

Met Ala Trp Asp Met Met Met Asn Trp
      180            185

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<210> 25

<211> 606
 <212> DNA
 <213> Hepatitis C virus

<220>
 <221> CDS
 <222> 1..603

<220>
 <221> mat_peptide
 <222> 1..600

<400> 25

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atg ttg ggt aag gtc atc gat acc ctt aca tgc ggc ttc gcc gac ctc
Met Leu Gly Lys Val Ile Asp Thr Leu Thr Cys Gly Phe Ala Asp Leu
 1             5             10             15

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gtg ggg tac att ccg ctc gtc ggc gcc ccc cta ggg ggc gct gcc agg      96
Val Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu Gly Gly Ala Ala Arg
      20                      25                      30

gcc ctg gcg cat ggc gtc cgg gtt ctg gag gac ggc gtg aac tat gca      144
Ala Leu Ala His Gly Val Arg Val Leu Glu Asp Gly Val Asn Tyr Ala
      35                      40                      45

aca ggg aat ttg ccc ggt tgc tct ttc tct atc ttc ctc ttg gct ttg      192
Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu
      50                      55                      60

ctg tcc tgt ctg acc gtt cca gct tcc gct tat gaa gtg cgc aac gtg      240
Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Tyr Glu Val Arg Asn Val
      65                      70                      75                      80

tcc ggg atg tac cat gtc acg aac gac tgc tcc aac tca agc att gtg      288
Ser Gly Met Tyr His Val Thr Asn Asp Cys Ser Asn Ser Ser Ile Val
      85                      90                      95

tat gag gca gcg gac atg atc atg cac acc ccc ggg tgc gtg ccc tgc      336
Tyr Glu Ala Ala Asp Met Ile Met His Thr Pro Gly Cys Val Pro Cys
      100                      105                      110

gtt cgg gag aac aac tct tcc cgc tgc tgg gta gcg ctc acc ccc acg      384
Val Arg Glu Asn Asn Ser Ser Arg Cys Trp Val Ala Leu Thr Pro Thr
      115                      120                      125

ctc gca gct agg aac gcc agc gtc ccc acc acg aca ata cga cgc cac      432
Leu Ala Ala Arg Asn Ala Ser Val Pro Thr Thr Thr Ile Arg Arg His
      130                      135                      140

gtc gat tcc cag ctg ttc acc atc tcg cct cgc cgg cat gag acg gtg      480
Val Asp Ser Gln Leu Phe Thr Ile Ser Pro Arg Arg His Glu Thr Val
      145                      150                      155                      160

cag gac tgc aat tgc tca atc tat ccc ggc cac ata acg ggt cac cgt      528
Gln Asp Cys Asn Cys Ser Ile Tyr Pro Gly His Ile Thr Gly His Arg
      165                      170                      175

atg gct tgg gat atg atg atg aac tgg tcg cct aca acg gcc ctg gtg      576
Met Ala Trp Asp Met Met Met Asn Trp Ser Pro Thr Thr Ala Leu Val
      180                      185                      190

gta tcg cag ctg ctc cgg atc ctc taatag      606
Val Ser Gln Leu Leu Arg Ile Leu
      195                      200

<210> 26

<211> 200
<212> PRT
<213> Hepatitis C virus

<400> 26

Met Leu Gly Lys Val Ile Asp Thr Leu Thr Cys Gly Phe Ala Asp Leu
  1                      5                      10                      15

Val Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu Gly Gly Ala Ala Arg
      20                      25                      30

Ala Leu Ala His Gly Val Arg Val Leu Glu Asp Gly Val Asn Tyr Ala

```

35 40 45
 Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu
 50 55 60
 Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Tyr Glu Val Arg Asn Val
 65 70 75 80
 Ser Gly Met Tyr His Val Thr Asn Asp Cys Ser Asn Ser Ser Ile Val
 85 90 95
 Tyr Glu Ala Ala Asp Met Ile Met His Thr Pro Gly Cys Val Pro Cys
 100 105 110
 Val Arg Glu Asn Asn Ser Ser Arg Cys Trp Val Ala Leu Thr Pro Thr
 115 120 125
 Leu Ala Ala Arg Asn Ala Ser Val Pro Thr Thr Thr Ile Arg Arg His
 130 135 140
 Val Asp Ser Gln Leu Phe Thr Ile Ser Pro Arg Arg His Glu Thr Val
 145 150 155 160
 Gln Asp Cys Asn Cys Ser Ile Tyr Pro Gly His Ile Thr Gly His Arg
 165 170 175
 Met Ala Trp Asp Met Met Met Asn Trp Ser Pro Thr Thr Ala Leu Val
 180 185 190
 Val Ser Gln Leu Leu Arg Ile Leu
 195 200

<210> 27

<211> 636

<212> DNA

<213> Hepatitis C virus

<220>

<221> CDS

<222> 1..633

<220>

<221> mat_peptide

<222> 1..630

<400> 27

atg ttg ggt aag gtc atc gat acc ctt aca tgc ggc ttc gcc gac ctc 48
 Met Leu Gly Lys Val Ile Asp Thr Leu Thr Cys Gly Phe Ala Asp Leu
 1 5 10 15
 gtg ggg tac att ccg ctc gtc ggc gcc ccc cta ggg ggc gct gcc agg 96
 Val Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu Gly Gly Ala Ala Arg
 20 25 30
 gcc ctg gcg cat ggc gtc cgg gtt ctg gag gac ggc gtg aac tat gca 144
 Ala Leu Ala His Gly Val Arg Val Leu Glu Asp Gly Val Asn Tyr Ala
 35 40 45
 aca ggg aat ttg ccc ggt tgc tct ttc tct atc ttc ctc ttg gct ttg 192
 Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu
 50 55 60

ctg tcc tgt ctg acc gtt cca gct tcc gct tat gaa gtg cgc aac gtg 240
 Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Tyr Glu Val Arg Asn Val
 65 70 75 80

tcc ggg atg tac cat gtc acg aac gac tgc tcc aac tca agc att gtg 288
 Ser Gly Met Tyr His Val Thr Asn Asp Cys Ser Asn Ser Ser Ile Val
 85 90 95

tat gag gca gcg gac atg atc atg cac acc ccc ggg tgc gtg ccc tgc 336
 Tyr Glu Ala Ala Asp Met Ile Met His Thr Pro Gly Cys Val Pro Cys
 100 105 110

gtt cgg gag aac aac tct tcc cgc tgc tgg gta gcg ctc acc ccc acg 384
 Val Arg Glu Asn Asn Ser Ser Arg Cys Trp Val Ala Leu Thr Pro Thr
 115 120 125

ctc gca gct agg aac gcc agc gtc ccc acc acg aca ata cga cgc cac 432
 Leu Ala Ala Arg Asn Ala Ser Val Pro Thr Thr Thr Ile Arg Arg His
 130 135 140

gtc gat tcc cag ctg ttc acc atc tcg cct cgc cgg cat gag acg gtg 480
 Val Asp Ser Gln Leu Phe Thr Ile Ser Pro Arg Arg His Glu Thr Val
 145 150 155 160

cag gac tgc aat tgc tca atc tat ccc ggc cac ata acg ggt cac cgt 528
 Gln Asp Cys Asn Cys Ser Ile Tyr Pro Gly His Ile Thr Gly His Arg
 165 170 175

atg gct tgg gat atg atg atg aac tgg tcg cct aca acg gcc ctg gtg 576
 Met Ala Trp Asp Met Met Met Asn Trp Ser Pro Thr Thr Ala Leu Val
 180 185 190

gta tcg cag ctg ctc cgg atc gtg atc gag ggc aga cac cat cac cac 624
 Val Ser Gln Leu Leu Arg Ile Val Ile Glu Gly Arg His His His His
 195 200 205

cat cac taatag 636
 His His
 210
 <210> 28

<211> 210
 <212> PRT
 <213> Hepatitis C virus

<400> 28

Met Leu Gly Lys Val Ile Asp Thr Leu Thr Cys Gly Phe Ala Asp Leu
 1 5 10 15

Val Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu Gly Gly Ala Ala Arg
 20 25 30

Ala Leu Ala His Gly Val Arg Val Leu Glu Asp Gly Val Asn Tyr Ala
 35 40 45

Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu
 50 55 60

Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Tyr Glu Val Arg Asn Val
 65 70 75 80

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<210> 29
<211> 630
<212> DNA
<213> Hepatitis C virus

<220>
<221> CDS
<222> 1..627

<220>
<221> mat_peptide
<222> 1..624
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<400> 29																
atg	ggg	aag	gtc	atc	gat	acc	ctt	acg	tgc	gga	ttc	gcc	gat	ctc	atg	48
Met	Gly	Lys	Val	Ile	Asp	Thr	Leu	Thr	Cys	Gly	Phe	Ala	Asp	Leu	Met	
1				5					10					15		
ggg	tac	atc	ccg	ctc	gtc	ggc	gct	ccc	gta	gga	ggc	gtc	gca	aga	gcc	96
Gly	Tyr	Ile	Pro	Leu	Val	Gly	Ala	Pro	Val	Gly	Gly	Val	Ala	Arg	Ala	
			20					25					30			
ctt	gcg	cat	ggc	gtg	agg	gcc	ctt	gaa	gac	ggg	ata	aat	ttc	gca	aca	144
Leu	Ala	His	Gly	Val	Arg	Ala	Leu	Glu	Asp	Gly	Ile	Asn	Phe	Ala	Thr	
		35					40					45				
ggg	aat	ttg	ccc	ggg	tgc	tcc	ttt	tct	att	ttc	ctt	ctc	gct	ctg	ttc	192
Gly	Asn	Leu	Pro	Gly	Cys	Ser	Phe	Ser	Ile	Phe	Leu	Leu	Ala	Leu	Phe	
	50					55					60					
tct	tgc	tta	att	cat	cca	gca	gct	agt	cta	gag	tgg	cgg	aat	acg	tct	240
Ser	Cys	Leu	Ile	His	Pro	Ala	Ala	Ser	Leu	Glu	Trp	Arg	Asn	Thr	Ser	
65					70					75				80		

ggc ctc tat gtc ctt acc aac gac tgt tcc aat agc agt att gtg tac 288
 Gly Leu Tyr Val Leu Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr
 85 90 95

gag gcc gat gac gtt att ctg cac aca ccc ggc tgc ata cct tgt gtc 336
 Glu Ala Asp Asp Val Ile Leu His Thr Pro Gly Cys Ile Pro Cys Val
 100 105 110

cag gac ggc aat aca tcc acg tgc tgg acc cca gtg aca cct aca gtg 384
 Gln Asp Gly Asn Thr Ser Thr Cys Trp Thr Pro Val Thr Pro Thr Val
 115 120 125

gca gtc aag tac gtc gga gca acc acc gct tgc ata cgc agt cat gtg 432
 Ala Val Lys Tyr Val Gly Ala Thr Thr Ala Ser Ile Arg Ser His Val
 130 135 140

gac cta tta gtg ggc gcg gcc acg atg tgc tct gcg ctc tac gtg ggt 480
 Asp Leu Leu Val Gly Ala Ala Thr Met Cys Ser Ala Leu Tyr Val Gly
 145 150 155 160

gac atg tgt ggg gct gtc ttc ctc gtg gga caa gcc ttc acg ttc aga 528
 Asp Met Cys Gly Ala Val Phe Leu Val Gly Gln Ala Phe Thr Phe Arg
 165 170 175

cct cgt cgc cat caa acg gtc cag acc tgt aac tgc tgc ctg tac cca 576
 Pro Arg Arg His Gln Thr Val Gln Thr Cys Asn Cys Ser Leu Tyr Pro
 180 185 190

ggc cat ctt tca gga cat cga atg gct tgg gat atg atg atg aac tgg 624
 Gly His Leu Ser Gly His Arg Met Ala Trp Asp Met Met Met Asn Trp
 195 200 205

taatag 630

<210> 30

<211> 208

<212> PRT

<213> Hepatitis C virus

<400> 30

Met Gly Lys Val Ile Asp Thr Leu Thr Cys Gly Phe Ala Asp Leu Met
 1 5 10 15

Gly Tyr Ile Pro Leu Val Gly Ala Pro Val Gly Gly Val Ala Arg Ala
 20 25 30

Leu Ala His Gly Val Arg Ala Leu Glu Asp Gly Ile Asn Phe Ala Thr
 35 40 45

Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu Phe
 50 55 60

Ser Cys Leu Ile His Pro Ala Ala Ser Leu Glu Trp Arg Asn Thr Ser
 65 70 75 80

Gly Leu Tyr Val Leu Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr
 85 90 95

Glu Ala Asp Asp Val Ile Leu His Thr Pro Gly Cys Ile Pro Cys Val
 100 105 110

Gln Asp Gly Asn Thr Ser Thr Cys Trp Thr Pro Val Thr Pro Thr Val
 115 120 125

Ala Val Lys Tyr Val Gly Ala Thr Thr Ala Ser Ile Arg Ser His Val
 130 135 140

Asp Leu Leu Val Gly Ala Ala Thr Met Cys Ser Ala Leu Tyr Val Gly
 145 150 155 160

Asp Met Cys Gly Ala Val Phe Leu Val Gly Gln Ala Phe Thr Phe Arg
 165 170 175

Pro Arg Arg His Gln Thr Val Gln Thr Cys Asn Cys Ser Leu Tyr Pro
 180 185 190

Gly His Leu Ser Gly His Arg Met Ala Trp Asp Met Met Met Asn Trp
 195 200 205

<210> 31

<211> 630

<212> DNA

<213> Hepatitis C virus

<220>

<221> CDS

<222> 1..627

<220>

<221> mat_peptide

<222> 1..624

<400> 31

atg ggt aag gtc atc gat acc cta acg tgc gga ttc gcc gat ctc atg	48
Met Gly Lys Val Ile Asp Thr Leu Thr Cys Gly Phe Ala Asp Leu Met	
1 5 10 15	
ggg tat atc ccg ctc gta ggc ggc ccc att ggg ggc gtc gca agg gct	96
Gly Tyr Ile Pro Leu Val Gly Gly Pro Ile Gly Gly Val Ala Arg Ala	
20 25 30	
ctc gca cac ggt gtg agg gtc ctt gag gac ggg gta aac tat gca aca	144
Leu Ala His Gly Val Arg Val Leu Glu Asp Gly Val Asn Tyr Ala Thr	
35 40 45	
ggg aat tta ccc ggt tgc tct ttc tct atc ttt att ctt gct ctt ctc	192
Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile Phe Ile Leu Ala Leu Leu	
50 55 60	
tcg tgt ctg acc gtt ccg gcc tct gca gtt ccc tac cga aat gcc tct	240
Ser Cys Leu Thr Val Pro Ala Ser Ala Val Pro Tyr Arg Asn Ala Ser	
65 70 75 80	
ggg att tat cat gtt acc aat gat tgc cca aac tct tcc ata gtc tat	288
Gly Ile Tyr His Val Thr Asn Asp Cys Pro Asn Ser Ser Ile Val Tyr	
85 90 95	
gag gca gat aac ctg atc cta cac gca cct ggt tgc gtg cct tgt gtc	336
Glu Ala Asp Asn Leu Ile Leu His Ala Pro Gly Cys Val Pro Cys Val	
100 105 110	
atg aca ggt aat gtg agt aga tgc tgg gtc caa att acc cct aca ctg	384

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Met Thr Gly Asn Val Ser Arg Cys Trp Val Gln Ile Thr Pro Thr Leu
    115                      120                      125

tca gcc ccg agc ctc gga gca gtc acg gct cct ctt cgg aga gcc gtt      432
Ser Ala Pro Ser Leu Gly Ala Val Thr Ala Pro Leu Arg Arg Ala Val
    130                      135                      140

gac tac cta gcg gga ggg gct gcc ctc tgc tcc gcg tta tac gta gga      480
Asp Tyr Leu Ala Gly Gly Ala Ala Leu Cys Ser Ala Leu Tyr Val Gly
    145                      150                      155

gac gcg tgt ggg gca cta ttc ttg gta ggc caa atg ttc acc tat agg      528
Asp Ala Cys Gly Ala Leu Phe Leu Val Gly Gln Met Phe Thr Tyr Arg
    165                      170                      175

cct cgc cag cac gct acg gtg cag aac tgc aac tgt tcc att tac agt      576
Pro Arg Gln His Ala Thr Val Gln Asn Cys Asn Cys Ser Ile Tyr Ser
    180                      185                      190

ggc cat gtt acc ggc cac cgg atg gca tgg gat atg atg atg aac tgg      624
Gly His Val Thr Gly His Arg Met Ala Trp Asp Met Met Met Asn Trp
    195                      200                      205

taatag                                                                630

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<210> 32
 <211> 208
 <212> PRT
 <213> Hepatitis C virus

<400> 32

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Met Gly Lys Val Ile Asp Thr Leu Thr Cys Gly Phe Ala Asp Leu Met
  1                      5                      10                      15

Gly Tyr Ile Pro Leu Val Gly Gly Pro Ile Gly Gly Val Ala Arg Ala
    20                      25                      30

Leu Ala His Gly Val Arg Val Leu Glu Asp Gly Val Asn Tyr Ala Thr
    35                      40                      45

Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile Phe Ile Leu Ala Leu Leu
    50                      55                      60

Ser Cys Leu Thr Val Pro Ala Ser Ala Val Pro Tyr Arg Asn Ala Ser
    65                      70                      75                      80

Gly Ile Tyr His Val Thr Asn Asp Cys Pro Asn Ser Ser Ile Val Tyr
    85                      90                      95

Glu Ala Asp Asn Leu Ile Leu His Ala Pro Gly Cys Val Pro Cys Val
    100                      105                      110

Met Thr Gly Asn Val Ser Arg Cys Trp Val Gln Ile Thr Pro Thr Leu
    115                      120                      125

Ser Ala Pro Ser Leu Gly Ala Val Thr Ala Pro Leu Arg Arg Ala Val
    130                      135                      140

Asp Tyr Leu Ala Gly Gly Ala Ala Leu Cys Ser Ala Leu Tyr Val Gly
    145                      150                      155                      160

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Asp Ala Cys Gly Ala Leu Phe Leu Val Gly Gln Met Phe Thr Tyr Arg
 165 170 175

Pro Arg Gln His Ala Thr Val Gln Asn Cys Asn Cys Ser Ile Tyr Ser
 180 185 190

Gly His Val Thr Gly His Arg Met Ala Trp Asp Met Met Met Asn Trp
 195 200 205

<210> 33

<211> 23

<212> DNA

<213> Hepatitis C virus

<400> 33

tgggatatga tgatgaactg gtc

23

<210> 34

<211> 30

<212> DNA

<213> Hepatitis C virus

<400> 34

ctattatggt ggtaagccac agagcaggag

30

<210> 35

<211> 1476

<212> DNA

<213> Hepatitis C virus

<220>

<221> CDS

<222> 1..1473

<220>

<221> mat_peptide

<222> 1..1470

<400> 35

tgg gat atg atg atg aac tgg tcg cct aca acg gcc ctg gtg gta tcg
 Trp Asp Met Met Met Asn Trp Ser Pro Thr Thr Ala Leu Val Val Ser
 1 5 10 15

48

cag ctg ctc cgg atc cca caa gct gtc gtg gac atg gtg gcg ggg gcc
 Gln Leu Leu Arg Ile Pro Gln Ala Val Val Asp Met Val Ala Gly Ala
 20 25 30

96

cat tgg gga gtc ctg gcg ggc ctc gcc tac tat tcc atg gtg ggg aac
 His Trp Gly Val Leu Ala Gly Leu Ala Tyr Tyr Ser Met Val Gly Asn
 35 40 45

144

tgg gct aag gtt ttg gtt gtg atg cta ctc ttt gcc ggc gtc gac ggg
 Trp Ala Lys Val Leu Val Val Met Leu Leu Phe Ala Gly Val Asp Gly
 50 55 60

192

cat acc cgc gtg tca gga ggg gca gca gcc tcc gat acc agg ggc ctt

240

His	Thr	Arg	Val	Ser	Gly	Gly	Ala	Ala	Ala	Ser	Asp	Thr	Arg	Gly	Leu	
65					70					75					80	
gtg	tcc	ctc	ttt	agc	ccc	ggg	tcg	gct	cag	aaa	atc	cag	ctc	gta	aac	288
Val	Ser	Leu	Phe	Ser	Pro	Gly	Ser	Ala	Gln	Lys	Ile	Gln	Leu	Val	Asn	
				85				90						95		
acc	aac	ggc	agt	tgg	cac	atc	aac	agg	act	gcc	ctg	aac	tgc	aac	gac	336
Thr	Asn	Gly	Ser	Trp	His	Ile	Asn	Arg	Thr	Ala	Leu	Asn	Cys	Asn	Asp	
			100					105					110			
tcc	ctc	caa	aca	ggg	ttc	ttt	gcc	gca	cta	ttc	tac	aaa	cac	aaa	ttc	384
Ser	Leu	Gln	Thr	Gly	Phe	Phe	Ala	Ala	Leu	Phe	Tyr	Lys	His	Lys	Phe	
		115					120					125				
aac	tcg	tct	gga	tgc	cca	gag	cgc	ttg	gcc	agc	tgt	cgc	tcc	atc	gac	432
Asn	Ser	Ser	Gly	Cys	Pro	Glu	Arg	Leu	Ala	Ser	Cys	Arg	Ser	Ile	Asp	
	130					135					140					
aag	ttc	gct	cag	ggg	tgg	ggt	ccc	ctc	act	tac	act	gag	cct	aac	agc	480
Lys	Phe	Ala	Gln	Gly	Trp	Gly	Pro	Leu	Thr	Tyr	Thr	Glu	Pro	Asn	Ser	
145				150						155					160	
tcg	gac	cag	agg	ccc	tac	tgc	tgg	cac	tac	gcg	cct	cga	ccg	tgt	ggt	528
Ser	Asp	Gln	Arg	Pro	Tyr	Cys	Trp	His	Tyr	Ala	Pro	Arg	Pro	Cys	Gly	
				165					170					175		
att	gta	ccc	gcg	tct	cag	gtg	tgc	ggt	cca	gtg	tat	tgc	ttc	acc	ccg	576
Ile	Val	Pro	Ala	Ser	Gln	Val	Cys	Gly	Pro	Val	Tyr	Cys	Phe	Thr	Pro	
			180					185					190			
agc	cct	gtt	gtg	gtg	ggg	acg	acc	gat	cgg	ttt	ggt	gtc	ccc	acg	tat	624
Ser	Pro	Val	Val	Val	Gly	Thr	Thr	Asp	Arg	Phe	Gly	Val	Pro	Thr	Tyr	
		195					200					205				
aac	tgg	ggg	gcg	aac	gac	tcg	gat	gtg	ctg	att	ctc	aac	aac	acg	cgg	672
Asn	Trp	Gly	Ala	Asn	Asp	Ser	Asp	Val	Leu	Ile	Leu	Asn	Asn	Thr	Arg	
	210					215					220					
ccg	ccg	cga	ggc	aac	tgg	ttc	ggc	tgt	aca	tgg	atg	aat	ggc	act	ggg	720
Pro	Pro	Arg	Gly	Asn	Trp	Phe	Gly	Cys	Thr	Trp	Met	Asn	Gly	Thr	Gly	
225				230						235				240		
ttc	acc	aag	acg	tgt	ggg	ggc	ccc	ccg	tgc	aac	atc	ggg	ggg	gcc	ggc	768
Phe	Thr	Lys	Thr	Cys	Gly	Gly	Pro	Pro	Cys	Asn	Ile	Gly	Gly	Ala	Gly	
				245					250					255		
aac	aac	acc	ttg	acc	tgc	ccc	act	gac	tgt	ttt	cgg	aag	cac	ccc	gag	816
Asn	Asn	Thr	Leu	Thr	Cys	Pro	Thr	Asp	Cys	Phe	Arg	Lys	His	Pro	Glu	
			260					265					270			
gcc	acc	tac	gcc	aga	tgc	ggt	tct	ggg	ccc	tgg	ctg	aca	cct	agg	tgt	864
Ala	Thr	Tyr	Ala	Arg	Cys	Gly	Ser	Gly	Pro	Trp	Leu	Thr	Pro	Arg	Cys	
		275					280						285			
atg	gtt	cat	tac	cca	tat	agg	ctc	tgg	cac	tac	ccc	tgc	act	gtc	aac	912
Met	Val	His	Tyr	Pro	Tyr	Arg	Leu	Trp	His	Tyr	Pro	Cys	Thr	Val	Asn	
	290					295					300					
ttc	acc	atc	ttc	aag	gtt	agg	atg	tac	gtg	ggg	ggc	gtg	gag	cac	agg	960
Phe	Thr	Ile	Phe	Lys	Val	Arg	Met	Tyr	Val	Gly	Gly	Val	Glu	His	Arg	
305					310					315					320	

ttc gaa gcc gca tgc aat tgg act cga gga gag cgt tgt gac ttg gag 1008
Phe Glu Ala Ala Cys Asn Trp Thr Arg Gly Glu Arg Cys Asp Leu Glu
325 330 335

gac agg gat aga tca gag ctt agc ccg ctg ctg ctg tct aca aca gag 1056
Asp Arg Asp Arg Ser Glu Leu Ser Pro Leu Leu Leu Ser Thr Thr Glu
340 345 350

tgg cag ata ctg ccc tgt tcc ttc acc acc ctg ccg gcc cta tcc acc 1104
 Trp Gln Ile Leu Pro Cys Ser Phe Thr Thr Leu Pro Ala Leu Ser Thr
 355 360 365

ggc ctg atc cac ctc cat cag aac atc gtg gac gtg caa tac ctg tac 1152
Gly Leu Ile His Leu His Gln Asn Ile Val Asp Val Gln Tyr Leu Tyr
370 375 380

ggt gta ggg tcg gcg gtt gtc tcc ctt gtc atc aaa tgg gag tat gtc 1200
Gly Val Gly Ser Ala Val Val Ser Leu Val Ile Lys Trp Glu Tyr Val
385 390 395 400

ctg	ttg	ctc	ttc	ctt	ctc	ctg	gca	gac	gcg	cgc	atc	tgc	gcc	tgc	tta	1248
Leu	Leu	Leu	Phe	Leu	Leu	Leu	Ala	Asp	Ala	Arg	Ile	Cys	Ala	Cys	Leu	
				405					410					415		

tgg atg atg ctg ctg ata gct caa gct gag gcc gcc tta gag aac ctg 1296
 Trp Met Met Leu Leu Ile Ala Gln Ala Glu Ala Ala Leu Glu Asn Leu
 420 425 430

gtg	gtc	ctc	aat	gcg	gcg	gcc	gtg	gcc	ggg	gcg	cat	ggc	act	ctt	tcc	1344
Val	Val	Leu	Asn	Ala	Ala	Ala	Val	Ala	Gly	Ala	His	Gly	Thr	Leu	Ser	
		435					440					445				

ttc ctt gtg ttc ttc tgt gct gcc tgg tac atc aag ggc agg ctg gtc 1392
Phe Leu Val Phe Phe Cys Ala Ala Trp Tyr Ile Lys Gly Arg Leu Val
450 455 460

cct ggt gcg gca tac gcc ttc tat ggc gtg tgg ccg ctg ctc ctg ctt 1440
Pro Gly Ala Ala Tyr Ala Phe Tyr Gly Val Trp Pro Leu Leu Leu Leu
465 470 475 480

ctg ctg gcc tta cca cca cga gct tat gcc tagtaa 1476
Leu Leu Ala Leu Pro Pro Arg Ala Tyr Ala
485 490

<210> 36

<211> 490

<212> PRT

<213> Hepatitis C virus

<400> 36

Trp Asp Met Met Met Asn Trp Ser Pro Thr Thr Ala Leu Val Val Ser
1 5 10 15

Gln Leu Leu Arg Ile Pro Gln Ala Val Val Asp Met Val Ala Gly Ala
20 25 30

His Trp Gly Val Leu Ala Gly Leu Ala Tyr Tyr Ser Met Val Gly Asn
35 40 45

Trp Ala Lys Val Leu Val Val Met Leu Leu Phe Ala Gly Val Asp Gly
50 55 60

His	Thr	Arg	Val	Ser	Gly	Gly	Ala	Ala	Ala	Ser	Asp	Thr	Arg	Gly	Leu	65	70	75	80
Val	Ser	Leu	Phe	Ser	Pro	Gly	Ser	Ala	Gln	Lys	Ile	Gln	Leu	Val	Asn	85	90	95	
Thr	Asn	Gly	Ser	Trp	His	Ile	Asn	Arg	Thr	Ala	Leu	Asn	Cys	Asn	Asp	100	105	110	
Ser	Leu	Gln	Thr	Gly	Phe	Phe	Ala	Ala	Leu	Phe	Tyr	Lys	His	Lys	Phe	115	120	125	
Asn	Ser	Ser	Gly	Cys	Pro	Glu	Arg	Leu	Ala	Ser	Cys	Arg	Ser	Ile	Asp	130	135	140	
Lys	Phe	Ala	Gln	Gly	Trp	Gly	Pro	Leu	Thr	Tyr	Thr	Glu	Pro	Asn	Ser	145	150	155	160
Ser	Asp	Gln	Arg	Pro	Tyr	Cys	Trp	His	Tyr	Ala	Pro	Arg	Pro	Cys	Gly	165	170	175	
Ile	Val	Pro	Ala	Ser	Gln	Val	Cys	Gly	Pro	Val	Tyr	Cys	Phe	Thr	Pro	180	185	190	
Ser	Pro	Val	Val	Val	Gly	Thr	Thr	Asp	Arg	Phe	Gly	Val	Pro	Thr	Tyr	195	200	205	
Asn	Trp	Gly	Ala	Asn	Asp	Ser	Asp	Val	Leu	Ile	Leu	Asn	Asn	Thr	Arg	210	215	220	
Pro	Pro	Arg	Gly	Asn	Trp	Phe	Gly	Cys	Thr	Trp	Met	Asn	Gly	Thr	Gly	225	230	235	240
Phe	Thr	Lys	Thr	Cys	Gly	Gly	Pro	Pro	Cys	Asn	Ile	Gly	Gly	Ala	Gly	245	250	255	
Asn	Asn	Thr	Leu	Thr	Cys	Pro	Thr	Asp	Cys	Phe	Arg	Lys	His	Pro	Glu	260	265	270	
Ala	Thr	Tyr	Ala	Arg	Cys	Gly	Ser	Gly	Pro	Trp	Leu	Thr	Pro	Arg	Cys	275	280	285	
Met	Val	His	Tyr	Pro	Tyr	Arg	Leu	Trp	His	Tyr	Pro	Cys	Thr	Val	Asn	290	295	300	
Phe	Thr	Ile	Phe	Lys	Val	Arg	Met	Tyr	Val	Gly	Gly	Val	Glu	His	Arg	305	310	315	320
Phe	Glu	Ala	Ala	Cys	Asn	Trp	Thr	Arg	Gly	Glu	Arg	Cys	Asp	Leu	Glu	325	330	335	
Asp	Arg	Asp	Arg	Ser	Glu	Leu	Ser	Pro	Leu	Leu	Leu	Ser	Thr	Thr	Glu	340	345	350	
Trp	Gln	Ile	Leu	Pro	Cys	Ser	Phe	Thr	Thr	Leu	Pro	Ala	Leu	Ser	Thr	355	360	365	
Gly	Leu	Ile	His	Leu	His	Gln	Asn	Ile	Val	Asp	Val	Gln	Tyr	Leu	Tyr	370	375	380	
Gly	Val	Gly	Ser	Ala	Val	Val	Ser	Leu	Val	Ile	Lys	Trp	Glu	Tyr	Val	385	390	395	400

Leu	Leu	Leu	Phe	Leu	Leu	Leu	Ala	Asp	Ala	Arg	Ile	Cys	Ala	Cys	Leu
			405					410						415	
Trp	Met	Met	Leu	Leu	Ile	Ala	Gln	Ala	Glu	Ala	Ala	Leu	Glu	Asn	Leu
			420					425					430		
Val	Val	Leu	Asn	Ala	Ala	Ala	Val	Ala	Gly	Ala	His	Gly	Thr	Leu	Ser
		435					440					445			
Phe	Leu	Val	Phe	Phe	Cys	Ala	Ala	Trp	Tyr	Ile	Lys	Gly	Arg	Leu	Val
	450					455					460				
Pro	Gly	Ala	Ala	Tyr	Ala	Phe	Tyr	Gly	Val	Trp	Pro	Leu	Leu	Leu	Leu
465					470					475					480
Leu	Leu	Ala	Leu	Pro	Pro	Arg	Ala	Tyr	Ala						
			485						490						

<210> 37

<211> 1021

<212> DNA

<213> Hepatitis C virus

<220>

<221> CDS

<222> 2..1018

<220>

<221> mat peptide

<222> 2..1015

<400> 37

g	atc	cca	caa	gct	gtc	gtg	gac	atg	gtg	gcg	ggg	gcc	cat	tgg	gga	46
Ile	Pro	Gln	Ala	Val	Val	Asp	Met	Val	Ala	Gly	Ala	His	Trp	Gly		
1				5						10				15		
gtc	ctg	gcg	ggc	ctc	gcc	tac	tat	tcc	atg	gtg	ggg	aac	tgg	gct	aag	94
Val	Leu	Ala	Gly	Leu	Ala	Tyr	Tyr	Ser	Met	Val	Gly	Asn	Trp	Ala	Lys	
				20					25					30		
gtt	ttg	gtt	gtg	atg	cta	ctc	ttt	gcc	ggc	gtc	gac	ggg	cat	acc	cgc	142
Val	Leu	Val	Val	Met	Leu	Leu	Phe	Ala	Gly	Val	Asp	Gly	His	Thr	Arg	
			35					40					45			
gtg	tca	gga	ggg	gca	gca	gcc	tcc	gat	acc	agg	ggc	ctt	gtg	tcc	ctc	190
Val	Ser	Gly	Gly	Ala	Ala	Ala	Ser	Asp	Thr	Arg	Gly	Leu	Val	Ser	Leu	
		50					55					60				
ttt	agc	ccc	ggg	tcg	gct	cag	aaa	atc	cag	ctc	gta	aac	acc	aac	ggc	238
Phe	Ser	Pro	Gly	Ser	Ala	Gln	Lys	Ile	Gln	Leu	Val	Asn	Thr	Asn	Gly	
	65					70					75					
agt	tgg	cac	atc	aac	agg	act	gcc	ctg	aac	tgc	aac	gac	tcc	ctc	caa	286
Ser	Trp	His	Ile	Asn	Arg	Thr	Ala	Leu	Asn	Cys	Asn	Asp	Ser	Leu	Gln	
80					85					90					95	
aca	ggg	ttc	ttt	gcc	gca	cta	ttc	tac	aaa	cac	aaa	ttc	aac	tcg	tct	334
Thr	Gly	Phe	Phe	Ala	Ala	Leu	Phe	Tyr	Lys	His	Lys	Phe	Asn	Ser	Ser	
				100					105					110		

gga tgc cca gag cgc ttg gcc agc tgt cgc tcc atc gac aag ttc gct 382
 Gly Cys Pro Glu Arg Leu Ala Ser Cys Arg Ser Ile Asp Lys Phe Ala
 115 120 125

 cag ggg tgg ggt ccc ctc act tac act gag cct aac agc tcg gac cag 430
 Gln Gly Trp Gly Pro Leu Thr Tyr Thr Glu Pro Asn Ser Ser Asp Gln
 130 135 140

 agg ccc tac tgc tgg cac tac gcg cct cga ccg tgt ggt att gta ccc 478
 Arg Pro Tyr Cys Trp His Tyr Ala Pro Arg Pro Cys Gly Ile Val Pro
 145 150 155

 gcg tct cag gtg tgc ggt cca gtg tat tgc ttc acc ccg agc cct gtt 526
 Ala Ser Gln Val Cys Gly Pro Val Tyr Cys Phe Thr Pro Ser Pro Val
 160 165 170 175
 gtg gtg ggg acg acc gat cgg ttt ggt gtc ccc acg tat aac tgg ggg 574
 Val Val Gly Thr Thr Asp Arg Phe Gly Val Pro Thr Tyr Asn Trp Gly
 180 185 190

 gcg aac gac tcg gat gtg ctg att ctc aac aac acg cgg ccg ccg cga 622
 Ala Asn Asp Ser Asp Val Leu Ile Leu Asn Asn Thr Arg Pro Pro Arg
 195 200 205

 ggc aac tgg ttc ggc tgt aca tgg atg aat ggc act ggg ttc acc aag 670
 Gly Asn Trp Phe Gly Cys Thr Trp Met Asn Gly Thr Gly Phe Thr Lys
 210 215 220

 acg tgt ggg ggc ccc ccg tgc aac atc ggg ggg gcc ggc aac aac acc 718
 Thr Cys Gly Gly Pro Pro Cys Asn Ile Gly Gly Ala Gly Asn Asn Thr
 225 230 235

 ttg acc tgc ccc act gac tgt ttt cgg aag cac ccc gag gcc acc tac 766
 Leu Thr Cys Pro Thr Asp Cys Phe Arg Lys His Pro Glu Ala Thr Tyr
 240 245 250 255

 gcc aga tgc ggt tct ggg ccc tgg ctg aca cct agg tgt atg gtt cat 814
 Ala Arg Cys Gly Ser Gly Pro Trp Leu Thr Pro Arg Cys Met Val His
 260 265 270

 tac cca tat agg ctc tgg cac tac ccc tgc act gtc aac ttc acc atc 862
 Tyr Pro Tyr Arg Leu Trp His Tyr Pro Cys Thr Val Asn Phe Thr Ile
 275 280 285

 ttc aag gtt agg atg tac gtg ggg ggc gtg gag cac agg ttc gaa gcc 910
 Phe Lys Val Arg Met Tyr Val Gly Gly Val Glu His Arg Phe Glu Ala
 290 295 300

 gca tgc aat tgg act cga gga gag cgt tgt gac ttg gag gac agg gat 958
 Ala Cys Asn Trp Thr Arg Gly Glu Arg Cys Asp Leu Glu Asp Arg Asp
 305 310 315

 aga tca gag ctt agc ccg ctg ctg ctg tct aca aca gag tgg cag agt 1006
 Arg Ser Glu Leu Ser Pro Leu Leu Leu Ser Thr Thr Glu Trp Gln Ser
 320 325 330 335

 ggc aga gct taatta 1021
 Gly Arg Ala

<210> 38

<211> 338

<212> PRT

<213> Hepatitis C virus

<400> 38

Ile Pro Gln Ala Val Val Asp Met Val Ala Gly Ala His Trp Gly Val
 1 5 10 15
 Leu Ala Gly Leu Ala Tyr Tyr Ser Met Val Gly Asn Trp Ala Lys Val
 20 25 30
 Leu Val Val Met Leu Leu Phe Ala Gly Val Asp Gly His Thr Arg Val
 35 40 45
 Ser Gly Gly Ala Ala Ala Ser Asp Thr Arg Gly Leu Val Ser Leu Phe
 50 55 60
 Ser Pro Gly Ser Ala Gln Lys Ile Gln Leu Val Asn Thr Asn Gly Ser
 65 70 75 80
 Trp His Ile Asn Arg Thr Ala Leu Asn Cys Asn Asp Ser Leu Gln Thr
 85 90 95
 Gly Phe Phe Ala Ala Leu Phe Tyr Lys His Lys Phe Asn Ser Ser Gly
 100 105 110
 Cys Pro Glu Arg Leu Ala Ser Cys Arg Ser Ile Asp Lys Phe Ala Gln
 115 120 125
 Gly Trp Gly Pro Leu Thr Tyr Thr Glu Pro Asn Ser Ser Asp Gln Arg
 130 135 140
 Pro Tyr Cys Trp His Tyr Ala Pro Arg Pro Cys Gly Ile Val Pro Ala
 145 150 155 160
 Ser Gln Val Cys Gly Pro Val Tyr Cys Phe Thr Pro Ser Pro Val Val
 165 170 175
 Val Gly Thr Thr Asp Arg Phe Gly Val Pro Thr Tyr Asn Trp Gly Ala
 180 185 190
 Asn Asp Ser Asp Val Leu Ile Leu Asn Asn Thr Arg Pro Pro Arg Gly
 195 200 205
 Asn Trp Phe Gly Cys Thr Trp Met Asn Gly Thr Gly Phe Thr Lys Thr
 210 215 220
 Cys Gly Gly Pro Pro Cys Asn Ile Gly Gly Ala Gly Asn Asn Thr Leu
 225 230 235 240
 Thr Cys Pro Thr Asp Cys Phe Arg Lys His Pro Glu Ala Thr Tyr Ala
 245 250 255
 Arg Cys Gly Ser Gly Pro Trp Leu Thr Pro Arg Cys Met Val His Tyr
 260 265 270
 Pro Tyr Arg Leu Trp His Tyr Pro Cys Thr Val Asn Phe Thr Ile Phe
 275 280 285
 Lys Val Arg Met Tyr Val Gly Gly Val Glu His Arg Phe Glu Ala Ala
 290 295 300
 Cys Asn Trp Thr Arg Gly Glu Arg Cys Asp Leu Glu Asp Arg Asp Arg
 305 310 315 320

Ser Glu Leu Ser Pro Leu Leu Leu Ser Thr Thr Glu Trp Gln Ser Gly
 325 330 335

Arg Ala

<210> 39

<211> 1034

<212> DNA

<213> Hepatitis C virus

<220>

<221> CDS

<222> 2..1032

<220>

<221> mat_peptide

<222> 2..1029

<400> 39

g atc cca caa gct gtc gtg gac atg gtg gcg ggg gcc cat tgg gga	46
Ile Pro Gln Ala Val Val Asp Met Val Ala Gly Ala His Trp Gly	
1 5 10 15	
gtc ctg gcg ggc ctc gcc tac tat tcc atg gtg ggg aac tgg gct aag	94
Val Leu Ala Gly Leu Ala Tyr Tyr Ser Met Val Gly Asn Trp Ala Lys	
20 25 30	
gtt ttg gtt gtg atg cta ctc ttt gcc ggc gtc gac ggg cat acc cgc	142
Val Leu Val Val Met Leu Leu Phe Ala Gly Val Asp Gly His Thr Arg	
35 40 45	
gtg tca gga ggg gca gca gcc tcc gat acc agg ggc ctt gtg tcc ctc	190
Val Ser Gly Gly Ala Ala Ala Ser Asp Thr Arg Gly Leu Val Ser Leu	
50 55 60	
ttt agc ccc ggg tcg gct cag aaa atc cag ctc gta aac acc aac ggc	238
Phe Ser Pro Gly Ser Ala Gln Lys Ile Gln Leu Val Asn Thr Asn Gly	
65 70 75	
agt tgg cac atc aac agg act gcc ctg aac tgc aac gac tcc ctc caa	286
Ser Trp His Ile Asn Arg Thr Ala Leu Asn Cys Asn Asp Ser Leu Gln	
80 85 90 95	
aca ggg ttc ttt gcc gca cta ttc tac aaa cac aaa ttc aac tcg tct	334
Thr Gly Phe Phe Ala Ala Leu Phe Tyr Lys His Lys Phe Asn Ser Ser	
100 105 110	
gga tgc cca gag cgc ttg gcc agc tgt cgc tcc atc gac aag ttc gct	382
Gly Cys Pro Glu Arg Leu Ala Ser Cys Arg Ser Ile Asp Lys Phe Ala	
115 120 125	
cag ggg tgg ggt ccc ctc act tac act gag cct aac agc tcg gac cag	430
Gln Gly Trp Gly Pro Leu Thr Tyr Thr Glu Pro Asn Ser Ser Asp Gln	
130 135 140	
agg ccc tac tgc tgg cac tac gcg cct cga ccg tgt ggt att gta ccc	478
Arg Pro Tyr Cys Trp His Tyr Ala Pro Arg Pro Cys Gly Ile Val Pro	
145 150 155	

gcg tct cag gtg tgc ggt cca gtg tat tgc ttc acc ccg agc cct gtt 526
 Ala Ser Gln Val Cys Gly Pro Val Tyr Cys Phe Thr Pro Ser Pro Val
 160 165 170 175

gtg gtg ggg acg acc gat cgg ttt ggt gtc ccc acg tat aac tgg ggg 574
 Val Val Gly Thr Thr Asp Arg Phe Gly Val Pro Thr Tyr Asn Trp Gly
 180 185 190

gcg aac gac tcg gat gtg ctg att ctc aac aac acg cgg ccg ccg cga 622
 Ala Asn Asp Ser Asp Val Leu Ile Leu Asn Asn Thr Arg Pro Pro Arg
 195 200 205

ggc aac tgg ttc ggc tgt aca tgg atg aat ggc act ggg ttc acc aag 670
 Gly Asn Trp Phe Gly Cys Thr Trp Met Asn Gly Thr Gly Phe Thr Lys
 210 215 220

acg tgt ggg ggc ccc ccg tgc aac atc ggg ggg gcc ggc aac aac acc 718
 Thr Cys Gly Gly Pro Pro Cys Asn Ile Gly Gly Ala Gly Asn Asn Thr
 225 230 235

ttg acc tgc ccc act gac tgt ttt cgg aag cac ccc gag gcc acc tac 766
 Leu Thr Cys Pro Thr Asp Cys Phe Arg Lys His Pro Glu Ala Thr Tyr
 240 245 250 255

gcc aga tgc ggt tct ggg ccc tgg ctg aca cct agg tgt atg gtt cat 814
 Ala Arg Cys Gly Ser Gly Pro Trp Leu Thr Pro Arg Cys Met Val His
 260 265 270

tac cca tat agg ctc tgg cac tac ccc tgc act gtc aac ttc acc atc 862
 Tyr Pro Tyr Arg Leu Trp His Tyr Pro Cys Thr Val Asn Phe Thr Ile
 275 280 285

ttc aag gtt agg atg tac gtg ggg ggc gtg gag cac agg ttc gaa gcc 910
 Phe Lys Val Arg Met Tyr Val Gly Gly Val Glu His Arg Phe Glu Ala
 290 295 300

gca tgc aat tgg act cga gga gag cgt tgt gac ttg gag gac agg gat 958
 Ala Cys Asn Trp Thr Arg Gly Glu Arg Cys Asp Leu Glu Asp Arg Asp
 305 310 315

aga tca gag ctt agc ccg ctg ctg ctg tct aca aca ggt gat cga ggg 1006
 Arg Ser Glu Leu Ser Pro Leu Leu Leu Ser Thr Thr Gly Asp Arg Gly
 320 325 330 335

cag aca cca tca cca cca tca cta at ag 1034
 Gln Thr Pro Ser Pro Pro Ser Leu
 340

<210> 40

<211> 343

<212> PRT

<213> Hepatitis C virus

<400> 40

Ile Pro Gln Ala Val Val Asp Met Val Ala Gly Ala His Trp Gly Val
 1 5 10 15

Leu Ala Gly Leu Ala Tyr Tyr Ser Met Val Gly Asn Trp Ala Lys Val
 20 25 30

Leu Val Val Met Leu Leu Phe Ala Gly Val Asp Gly His Thr Arg Val

35	40	45
Ser Gly Gly Ala Ala Ala	Ser Asp Thr Arg Gly	Leu Val Ser Leu Phe
50	55	60
Ser Pro Gly Ser Ala Gln Lys Ile Gln Leu Val Asn Thr Asn Gly Ser		
65	70	75 80
Trp His Ile Asn Arg Thr Ala Leu Asn Cys Asn Asp Ser Leu Gln Thr		
	85	90 95
Gly Phe Phe Ala Ala Leu Phe Tyr Lys His Lys Phe Asn Ser Ser Gly		
	100	105 110
Cys Pro Glu Arg Leu Ala Ser Cys Arg Ser Ile Asp Lys Phe Ala Gln		
	115	120 125
Gly Trp Gly Pro Leu Thr Tyr Thr Glu Pro Asn Ser Ser Asp Gln Arg		
	130	135 140
Pro Tyr Cys Trp His Tyr Ala Pro Arg Pro Cys Gly Ile Val Pro Ala		
145	150	155 160
Ser Gln Val Cys Gly Pro Val Tyr Cys Phe Thr Pro Ser Pro Val Val		
	165	170 175
Val Gly Thr Thr Asp Arg Phe Gly Val Pro Thr Tyr Asn Trp Gly Ala		
	180	185 190
Asn Asp Ser Asp Val Leu Ile Leu Asn Asn Thr Arg Pro Pro Arg Gly		
	195	200 205
Asn Trp Phe Gly Cys Thr Trp Met Asn Gly Thr Gly Phe Thr Lys Thr		
	210	215 220
Cys Gly Gly Pro Pro Cys Asn Ile Gly Gly Ala Gly Asn Asn Thr Leu		
225	230	235 240
Thr Cys Pro Thr Asp Cys Phe Arg Lys His Pro Glu Ala Thr Tyr Ala		
	245	250 255
Arg Cys Gly Ser Gly Pro Trp Leu Thr Pro Arg Cys Met Val His Tyr		
	260	265 270
Pro Tyr Arg Leu Trp His Tyr Pro Cys Thr Val Asn Phe Thr Ile Phe		
	275	280 285
Lys Val Arg Met Tyr Val Gly Gly Val Glu His Arg Phe Glu Ala Ala		
	290	295 300
Cys Asn Trp Thr Arg Gly Glu Arg Cys Asp Leu Glu Asp Arg Asp Arg		
305	310	315 320
Ser Glu Leu Ser Pro Leu Leu Leu Ser Thr Thr Gly Asp Arg Gly Gln		
	325	330 335
Thr Pro Ser Pro Pro Ser Leu		
	340	

<210> 41

<211> 945

<212> DNA

<213> Hepatitis C virus

<220>
 <221> CDS
 <222> 1..942

<220>
 <221> mat_peptide
 <222> 1..939

<400> 41

atg gtg ggg aac tgg gct aag gtt ttg gtt gtg atg cta ctc ttt gcc	48
Met Val Gly Asn Trp Ala Lys Val Leu Val Val Met Leu Leu Phe Ala	
1 5 10 15	
ggc gtc gac ggg cat acc cgc gtg tca gga ggg gca gca gcc tcc gat	96
Gly Val Asp Gly His Thr Arg Val Ser Gly Gly Ala Ala Ala Ser Asp	
20 25 30	
acc agg ggc ctt gtg tcc ctc ttt agc ccc ggg tcg gct cag aaa atc	144
Thr Arg Gly Leu Val Ser Leu Phe Ser Pro Gly Ser Ala Gln Lys Ile	
35 40 45	
cag ctc gta aac acc aac ggc agt tgg cac atc aac agg act gcc ctg	192
Gln Leu Val Asn Thr Asn Gly Ser Trp His Ile Asn Arg Thr Ala Leu	
50 55 60	
aac tgc aac gac tcc ctc caa aca ggg ttc ttt gcc gca cta ttc tac	240
Asn Cys Asn Asp Ser Leu Gln Thr Gly Phe Phe Ala Ala Leu Phe Tyr	
65 70 75 80	
aaa cac aaa ttc aac tcg tct gga tgc cca gag cgc ttg gcc agc tgt	288
Lys His Lys Phe Asn Ser Ser Gly Cys Pro Glu Arg Leu Ala Ser Cys	
85 90 95	
cgc tcc atc gac aag ttc gct cag ggg tgg ggt ccc ctc act tac act	336
Arg Ser Ile Asp Lys Phe Ala Gln Gly Trp Gly Pro Leu Thr Tyr Thr	
100 105 110	
gag cct aac agc tcg gac cag agg ccc tac tgc tgg cac tac gcg cct	384
Glu Pro Asn Ser Ser Asp Gln Arg Pro Tyr Cys Trp His Tyr Ala Pro	
115 120 125	
cga ccg tgt ggt att gta ccc gcg tct cag gtg tgc ggt cca gtg tat	432
Arg Pro Cys Gly Ile Val Pro Ala Ser Gln Val Cys Gly Pro Val Tyr	
130 135 140	
tgc ttc acc ccg agc cct gtt gtg gtg ggg acg acc gat cgg ttt ggt	480
Cys Phe Thr Pro Ser Pro Val Val Val Gly Thr Thr Asp Arg Phe Gly	
145 150 155 160	
gtc ccc acg tat aac tgg ggg gcg aac gac tcg gat gtg ctg att ctc	528
Val Pro Thr Tyr Asn Trp Gly Ala Asn Asp Ser Asp Val Leu Ile Leu	
165 170 175	
aac aac acg ccg ccg ccg cga ggc aac tgg ttc ggc tgt aca tgg atg	576
Asn Asn Thr Arg Pro Pro Arg Gly Asn Trp Phe Gly Cys Thr Trp Met	
180 185 190	
aat ggc act ggg ttc acc aag acg tgt ggg ggc ccc ccg tgc aac atc	624
Asn Gly Thr Gly Phe Thr Lys Thr Cys Gly Gly Pro Pro Cys Asn Ile	
195 200 205	
ggg ggg gcc ggc aac aac acc ttg acc tgc ccc act gac tgt ttt cgg	672

Gly Gly Ala Gly Asn Asn Thr Leu Thr Cys Pro Thr Asp Cys Phe Arg
 210 215 220
 aag cac ccc gag gcc acc tac gcc aga tgc ggt tct ggg ccc tgg ctg 720
 Lys His Pro Glu Ala Thr Tyr Ala Arg Cys Gly Ser Gly Pro Trp Leu
 225 230 235 240
 aca cct agg tgt atg gtt cat tac cca tat agg ctc tgg cac tac ccc 768
 Thr Pro Arg Cys Met Val His Tyr Pro Tyr Arg Leu Trp His Tyr Pro
 245 250 255
 tgc act gtc aac ttc acc atc ttc aag gtt agg atg tac gtg ggg ggc 816
 Cys Thr Val Asn Phe Thr Ile Phe Lys Val Arg Met Tyr Val Gly Gly
 260 265 270
 gtg gag cac agg ttc gaa gcc gca tgc aat tgg act cga gga gag cgt 864
 Val Glu His Arg Phe Glu Ala Ala Cys Asn Trp Thr Arg Gly Glu Arg
 275 280 285
 tgt gac ttg gag gac agg gat aga tca gag ctt agc ccg ctg ctg ctg 912
 Cys Asp Leu Glu Asp Arg Asp Arg Ser Glu Leu Ser Pro Leu Leu Leu
 290 295 300
 tct aca aca gag tgg cag agc tta att aat tag 945
 Ser Thr Thr Glu Trp Gln Ser Leu Ile Asn
 305 310
 <210> 42
 <211> 314
 <212> PRT
 <213> Hepatitis C virus
 <400> 42
 Met Val Gly Asn Trp Ala Lys Val Leu Val Val Met Leu Leu Phe Ala
 1 5 10 15
 Gly Val Asp Gly His Thr Arg Val Ser Gly Gly Ala Ala Ala Ser Asp
 20 25 30
 Thr Arg Gly Leu Val Ser Leu Phe Ser Pro Gly Ser Ala Gln Lys Ile
 35 40 45
 Gln Leu Val Asn Thr Asn Gly Ser Trp His Ile Asn Arg Thr Ala Leu
 50 55 60
 Asn Cys Asn Asp Ser Leu Gln Thr Gly Phe Phe Ala Ala Leu Phe Tyr
 65 70 75 80
 Lys His Lys Phe Asn Ser Ser Gly Cys Pro Glu Arg Leu Ala Ser Cys
 85 90 95
 Arg Ser Ile Asp Lys Phe Ala Gln Gly Trp Gly Pro Leu Thr Tyr Thr
 100 105 110
 Glu Pro Asn Ser Ser Asp Gln Arg Pro Tyr Cys Trp His Tyr Ala Pro
 115 120 125
 Arg Pro Cys Gly Ile Val Pro Ala Ser Gln Val Cys Gly Pro Val Tyr
 130 135 140
 Cys Phe Thr Pro Ser Pro Val Val Val Gly Thr Thr Asp Arg Phe Gly
 145 150 155 160

Val Pro Thr Tyr Asn Trp Gly Ala Asn Asp Ser Asp Val Leu Ile Leu
165 170 175

Asn Asn Thr Arg Pro Pro Arg Gly Asn Trp Phe Gly Cys Thr Trp Met
180 185 190

Asn Gly Thr Gly Phe Thr Lys Thr Cys Gly Gly Pro Pro Cys Asn Ile
195 200 205

Gly Gly Ala Gly Asn Asn Thr Leu Thr Cys Pro Thr Asp Cys Phe Arg
210 215 220

Lys His Pro Glu Ala Thr Tyr Ala Arg Cys Gly Ser Gly Pro Trp Leu
225 230 235 240

Thr Pro Arg Cys Met Val His Tyr Pro Tyr Arg Leu Trp His Tyr Pro
245 250 255

Cys Thr Val Asn Phe Thr Ile Phe Lys Val Arg Met Tyr Val Gly Gly
260 265 270

Val Glu His Arg Phe Glu Ala Ala Cys Asn Trp Thr Arg Gly Glu Arg
275 280 285

Cys Asp Leu Glu Asp Arg Asp Arg Ser Glu Leu Ser Pro Leu Leu Leu
290 295 300

Ser Thr Thr Glu Trp Gln Ser Leu Ile Asn
305 310

<210> 43

<211> 961

<212> DNA

<213> Hepatitis C virus

<220>

<221> CDS

<222> 1..958

<220>

<221> mat_peptide

<222> 1..955

<400> 43

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Met Val Gly Asn Trp Ala Lys Val Leu Val Val Met Leu Leu Phe Ala	
1 5 10 15	
ggc gtc gac ggg cat acc cgc gtg tca gga ggg gca gca gcc tcc gat	96
Gly Val Asp Gly His Thr Arg Val Ser Gly Gly Ala Ala Ala Ser Asp	
20 25 30	
acc agg ggc ctt gtg tcc ctc ttt agc ccc ggg tcg gct cag aaa atc	144
Thr Arg Gly Leu Val Ser Leu Phe Ser Pro Gly Ser Ala Gln Lys Ile	
35 40 45	
cag ctc gta aac acc aac ggc agt tgg cac atc aac agg act gcc ctg	192
Gln Leu Val Asn Thr Asn Gly Ser Trp His Ile Asn Arg Thr Ala Leu	
50 55 60	

aac tgc aac gac tcc ctc caa aca ggg ttc ttt gcc gca cta ttc tac	240
Asn Cys Asn Asp Ser Leu Gln Thr Gly Phe Phe Ala Ala Leu Phe Tyr	
65 70 75 80	
aaa cac aaa ttc aac tcg tct gga tgc cca gag cgc ttg gcc agc tgt	288
Lys His Lys Phe Asn Ser Ser Gly Cys Pro Glu Arg Leu Ala Ser Cys	
85 90 95	
cgc tcc atc gac aag ttc gct cag ggg tgg ggt ccc ctc act tac act	336
Arg Ser Ile Asp Lys Phe Ala Gln Gly Trp Gly Pro Leu Thr Tyr Thr	
100 105 110	
gag cct aac agc tcg gac cag agg ccc tac tgc tgg cac tac gcg cct	384
Glu Pro Asn Ser Ser Asp Gln Arg Pro Tyr Cys Trp His Tyr Ala Pro	
115 120 125	
cga ccg tgt ggt att gta ccc gcg tct cag gtg tgc ggt cca gtg tat	432
Arg Pro Cys Gly Ile Val Pro Ala Ser Gln Val Cys Gly Pro Val Tyr	
130 135 140	
tgc ttc acc ccg agc cct gtt gtg gtg ggg acg acc gat cgg ttt ggt	480
Cys Phe Thr Pro Ser Pro Val Val Val Gly Thr Thr Asp Arg Phe Gly	
145 150 155 160	
gtc ccc acg tat aac tgg ggg gcg aac gac tcg gat gtg ctg att ctc	528
Val Pro Thr Tyr Asn Trp Gly Ala Asn Asp Ser Asp Val Leu Ile Leu	
165 170 175	
aac aac acg cgg ccg ccg cga ggc aac tgg ttc ggc tgt aca tgg atg	576
Asn Asn Thr Arg Pro Pro Arg Gly Asn Trp Phe Gly Cys Thr Trp Met	
180 185 190	
aat ggc act ggg ttc acc aag acg tgt ggg ggc ccc ccg tgc aac atc	624
Asn Gly Thr Gly Phe Thr Lys Thr Cys Gly Gly Pro Cys Asn Ile	
195 200 205	
ggg ggg gcc ggc aac aac acc ttg acc tgc ccc act gac tgt ttt cgg	672
Gly Gly Ala Gly Asn Asn Thr Leu Thr Cys Pro Thr Asp Cys Phe Arg	
210 215 220	
aag cac ccc gag gcc acc tac gcc aga tgc ggt tct ggg ccc tgg ctg	720
Lys His Pro Glu Ala Thr Tyr Ala Arg Cys Gly Ser Gly Pro Trp Leu	
225 230 235 240	
aca cct agg tgt atg gtt cat tac cca tat agg ctc tgg cac tac ccc	768
Thr Pro Arg Cys Met Val His Tyr Pro Tyr Arg Leu Trp His Tyr Pro	
245 250 255	
tgc act gtc aac ttc acc atc ttc aag gtt agg atg tac gtg ggg ggc	816
Cys Thr Val Asn Phe Thr Ile Phe Lys Val Arg Met Tyr Val Gly Gly	
260 265 270	
gtg gag cac agg ttc gaa gcc gca tgc aat tgg act cga gga gag cgt	864
Val Glu His Arg Phe Glu Ala Ala Cys Asn Trp Thr Arg Gly Glu Arg	
275 280 285	
tgt gac ttg gag gac agg gat aga tca gag ctt agc ccg ctg ctg ctg	912
Cys Asp Leu Glu Asp Arg Asp Arg Ser Glu Leu Ser Pro Leu Leu Leu	
290 295 300	
tct aca aca ggt gat cga ggg cag aca cca tca cca cca tca cta a	958
Ser Thr Thr Gly Asp Arg Gly Gln Thr Pro Ser Pro Pro Ser Leu	
305 310 315	

tag

961

<210> 44

<211> 319

<212> PRT

<213> Hepatitis C virus

<400> 44

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Met Val Gly Asn Trp Ala Lys Val Leu Val Val Met Leu Leu Phe Ala
 1           5           10           15

Gly Val Asp Gly His Thr Arg Val Ser Gly Gly Ala Ala Ala Ser Asp
          20           25           30

Thr Arg Gly Leu Val Ser Leu Phe Ser Pro Gly Ser Ala Gln Lys Ile
          35           40           45

Gln Leu Val Asn Thr Asn Gly Ser Trp His Ile Asn Arg Thr Ala Leu
 50           55           60

Asn Cys Asn Asp Ser Leu Gln Thr Gly Phe Phe Ala Ala Leu Phe Tyr
 65           70           75           80

Lys His Lys Phe Asn Ser Ser Gly Cys Pro Glu Arg Leu Ala Ser Cys
          85           90           95

Arg Ser Ile Asp Lys Phe Ala Gln Gly Trp Gly Pro Leu Thr Tyr Thr
          100          105          110

Glu Pro Asn Ser Ser Asp Gln Arg Pro Tyr Cys Trp His Tyr Ala Pro
          115          120          125

Arg Pro Cys Gly Ile Val Pro Ala Ser Gln Val Cys Gly Pro Val Tyr
          130          135          140

Cys Phe Thr Pro Ser Pro Val Val Val Gly Thr Thr Asp Arg Phe Gly
          145          150          155          160

Val Pro Thr Tyr Asn Trp Gly Ala Asn Asp Ser Asp Val Leu Ile Leu
          165          170          175

Asn Asn Thr Arg Pro Pro Arg Gly Asn Trp Phe Gly Cys Thr Trp Met
          180          185          190

Asn Gly Thr Gly Phe Thr Lys Thr Cys Gly Gly Pro Pro Cys Asn Ile
          195          200          205

Gly Gly Ala Gly Asn Asn Thr Leu Thr Cys Pro Thr Asp Cys Phe Arg
          210          215          220

Lys His Pro Glu Ala Thr Tyr Ala Arg Cys Gly Ser Gly Pro Trp Leu
          225          230          235          240

Thr Pro Arg Cys Met Val His Tyr Pro Tyr Arg Leu Trp His Tyr Pro
          245          250          255

Cys Thr Val Asn Phe Thr Ile Phe Lys Val Arg Met Tyr Val Gly Gly
          260          265          270

Val Glu His Arg Phe Glu Ala Ala Cys Asn Trp Thr Arg Gly Glu Arg

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                275                280                285
Cys Asp Leu Glu Asp Arg Asp Arg Ser Glu Leu Ser Pro Leu Leu Leu
 290                295                300

Ser Thr Thr Gly Asp Arg Gly Gln Thr Pro Ser Pro Pro Ser Leu
305                310                315

<210> 45

<211> 1395
<212> DNA
<213> Hepatitis C virus

<220>
<221> CDS
<222> 1..1392

<220>
<221> mat_peptide
<222> 1..1389

<400> 45
atg gtg gcg ggg gcc cat tgg gga gtc ctg gcg ggc ctc gcc tac tat      48
Met Val Ala Gly Ala His Trp Gly Val Leu Ala Gly Leu Ala Tyr Tyr
  1                5                10                15

tcc atg gtg ggg aac tgg gct aag gtt ttg gtt gtg atg cta ctc ttt      96
Ser Met Val Gly Asn Trp Ala Lys Val Leu Val Val Met Leu Leu Phe
          20                25                30

gcc ggc gtc gac ggg cat acc cgc gtg tca gga ggg gca gca gcc tcc      144
Ala Gly Val Asp Gly His Thr Arg Val Ser Gly Gly Ala Ala Ala Ser
          35                40                45

gat acc agg ggc ctt gtg tcc ctc ttt agc ccc ggg tcg gct cag aaa      192
Asp Thr Arg Gly Leu Val Ser Leu Phe Ser Pro Gly Ser Ala Gln Lys
          50                55                60

atc cag ctc gta aac acc aac ggc agt tgg cac atc aac agg act gcc      240
Ile Gln Leu Val Asn Thr Asn Gly Ser Trp His Ile Asn Arg Thr Ala
          65                70                75                80

ctg aac tgc aac gac tcc ctc caa aca ggg ttc ttt gcc gca cta ttc      288
Leu Asn Cys Asn Asp Ser Leu Gln Thr Gly Phe Phe Ala Ala Leu Phe
          85                90                95

tac aaa cac aaa ttc aac tcg tct gga tgc cca gag cgc ttg gcc agc      336
Tyr Lys His Lys Phe Asn Ser Ser Gly Cys Pro Glu Arg Leu Ala Ser
          100                105                110

tgt cgc tcc atc gac aag ttc gct cag ggg tgg ggt ccc ctc act tac      384
Cys Arg Ser Ile Asp Lys Phe Ala Gln Gly Trp Gly Pro Leu Thr Tyr
          115                120                125

act gag cct aac agc tcg gac cag agg ccc tac tgc tgg cac tac gcg      432
Thr Glu Pro Asn Ser Ser Asp Gln Arg Pro Tyr Cys Trp His Tyr Ala
          130                135                140

cct cga ccg tgt ggt att gta ccc gcg tct cag gtg tgc ggt cca gtg      480
Pro Arg Pro Cys Gly Ile Val Pro Ala Ser Gln Val Cys Gly Pro Val
          145                150                155                160

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tat tgc ttc acc ccg agc cct gtt gtg gtg ggg acg acc gat cgg ttt Tyr Cys Phe Thr Pro Ser Pro Val Val Val Gly Thr Thr Asp Arg Phe 165 170 175	528
ggg gtc ccc acg tat aac tgg ggg gcg aac gac tcg gat gtg ctg att Gly Val Pro Thr Tyr Asn Trp Gly Ala Asn Asp Ser Asp Val Leu Ile 180 185 190	576
ctc aac aac acg cgg ccg ccg cga ggc aac tgg ttc ggc tgt aca tgg Leu Asn Asn Thr Arg Pro Pro Arg Gly Asn Trp Phe Gly Cys Thr Trp 195 200 205	624
atg aat ggc act ggg ttc acc aag acg tgt ggg ggc ccc ccg tgc aac Met Asn Gly Thr Gly Phe Thr Lys Thr Cys Gly Gly Pro Pro Cys Asn 210 215 220	672
atc ggg ggg gcc ggc aac aac acc ttg acc tgc ccc act gac tgt ttt Ile Gly Gly Ala Gly Asn Asn Thr Leu Thr Cys Pro Thr Asp Cys Phe 225 230 235 240	720
cgg aag cac ccc gag gcc acc tac gcc aga tgc ggt tct ggg ccc tgg Arg Lys His Pro Glu Ala Thr Tyr Ala Arg Cys Gly Ser Gly Pro Trp 245 250 255	768
ctg aca cct agg tgt atg gtt cat tac cca tat agg ctc tgg cac tac Leu Thr Pro Arg Cys Met Val His Tyr Pro Tyr Arg Leu Trp His Tyr 260 265 270	816
ccc tgc act gtc aac ttc acc atc ttc aag gtt agg atg tac gtg ggg Pro Cys Thr Val Asn Phe Thr Ile Phe Lys Val Arg Met Tyr Val Gly 275 280 285	864
ggc gtg gag cac agg ttc gaa gcc gca tgc aat tgg act cga gga gag Gly Val Glu His Arg Phe Glu Ala Ala Cys Asn Trp Thr Arg Gly Glu 290 295 300	912
cgt tgt gac ttg gag gac agg gat aga tca gag ctt agc ccg ctg ctg Arg Cys Asp Leu Glu Asp Arg Asp Arg Ser Glu Leu Ser Pro Leu Leu 305 310 315 320	960
ctg tct aca aca gag tgg cag ata ctg ccc tgt tcc ttc acc acc ctg Leu Ser Thr Thr Glu Trp Gln Ile Leu Pro Cys Ser Phe Thr Thr Leu 325 330 335	1008
ccg gcc cta tcc acc ggc ctg atc cac ctc cat cag aac atc gtg gac Pro Ala Leu Ser Thr Gly Leu Ile His Leu His Gln Asn Ile Val Asp 340 345 350	1056
gtg caa tac ctg tac ggt gta ggg tcg gcg gtt gtc tcc ctt gtc atc Val Gln Tyr Leu Tyr Gly Val Gly Ser Ala Val Val Ser Leu Val Ile 355 360 365	1104
aaa tgg gag tat gtc ctg ttg ctc ttc ctt ctc ctg gca gac gcg cgc Lys Trp Glu Tyr Val Leu Leu Leu Phe Leu Leu Leu Ala Asp Ala Arg 370 375 380	1152
atc tgc gcc tgc tta tgg atg atg ctg ctg ata gct caa gct gag gcc Ile Cys Ala Cys Leu Trp Met Met Leu Leu Ile Ala Gln Ala Glu Ala 385 390 395 400	1200
gcc tta gag aac ctg gtg gtc ctc aat gcg gcg gcc gtg gcc ggg gcg Ala Leu Glu Asn Leu Val Val Leu Asn Ala Ala Ala Val Ala Gly Ala	1248

405 410 415
 cat ggc act ctt tcc ttc ctt gtg ttc ttc tgt gct gcc tgg tac atc 1296
 His Gly Thr Leu Ser Phe Leu Val Phe Phe Cys Ala Ala Trp Tyr Ile
 420 425 430
 aag ggc agg ctg gtc cct ggt gcg gca tac gcc ttc tat ggc gtg tgg 1344
 Lys Gly Arg Leu Val Pro Gly Ala Ala Tyr Ala Phe Tyr Gly Val Trp
 435 440 445
 ccg ctg ctc ctg ctt ctg ctg gcc tta cca cca cga gct tat gcc tagtaa 1395
 Pro Leu Leu Leu Leu Leu Leu Ala Leu Pro Pro Arg Ala Tyr Ala
 450 455 460
 <210> 46
 <211> 463
 <212> PRT
 <213> Hepatitis C virus
 <400> 46
 Met Val Ala Gly Ala His Trp Gly Val Leu Ala Gly Leu Ala Tyr Tyr
 1 5 10 15
 Ser Met Val Gly Asn Trp Ala Lys Val Leu Val Val Met Leu Leu Phe
 20 25 30
 Ala Gly Val Asp Gly His Thr Arg Val Ser Gly Gly Ala Ala Ala Ser
 35 40 45
 Asp Thr Arg Gly Leu Val Ser Leu Phe Ser Pro Gly Ser Ala Gln Lys
 50 55 60
 Ile Gln Leu Val Asn Thr Asn Gly Ser Trp His Ile Asn Arg Thr Ala
 65 70 75 80
 Leu Asn Cys Asn Asp Ser Leu Gln Thr Gly Phe Phe Ala Ala Leu Phe
 85 90 95
 Tyr Lys His Lys Phe Asn Ser Ser Gly Cys Pro Glu Arg Leu Ala Ser
 100 105 110
 Cys Arg Ser Ile Asp Lys Phe Ala Gln Gly Trp Gly Pro Leu Thr Tyr
 115 120 125
 Thr Glu Pro Asn Ser Ser Asp Gln Arg Pro Tyr Cys Trp His Tyr Ala
 130 135 140
 Pro Arg Pro Cys Gly Ile Val Pro Ala Ser Gln Val Cys Gly Pro Val
 145 150 155 160
 Tyr Cys Phe Thr Pro Ser Pro Val Val Val Gly Thr Thr Asp Arg Phe
 165 170 175
 Gly Val Pro Thr Tyr Asn Trp Gly Ala Asn Asp Ser Asp Val Leu Ile
 180 185 190
 Leu Asn Asn Thr Arg Pro Pro Arg Gly Asn Trp Phe Gly Cys Thr Trp
 195 200 205
 Met Asn Gly Thr Gly Phe Thr Lys Thr Cys Gly Gly Pro Pro Cys Asn
 210 215 220

Ile Gly Gly Ala Gly Asn Asn Thr Leu Thr Cys Pro Thr Asp Cys Phe
 225 230 235 240
 Arg Lys His Pro Glu Ala Thr Tyr Ala Arg Cys Gly Ser Gly Pro Trp
 245 250 255
 Leu Thr Pro Arg Cys Met Val His Tyr Pro Tyr Arg Leu Trp His Tyr
 260 265 270
 Pro Cys Thr Val Asn Phe Thr Ile Phe Lys Val Arg Met Tyr Val Gly
 275 280 285
 Gly Val Glu His Arg Phe Glu Ala Ala Cys Asn Trp Thr Arg Gly Glu
 290 295 300
 Arg Cys Asp Leu Glu Asp Arg Asp Arg Ser Glu Leu Ser Pro Leu Leu
 305 310 315 320
 Leu Ser Thr Thr Glu Trp Gln Ile Leu Pro Cys Ser Phe Thr Thr Leu
 325 330 335
 Pro Ala Leu Ser Thr Gly Leu Ile His Leu His Gln Asn Ile Val Asp
 340 345 350
 Val Gln Tyr Leu Tyr Gly Val Gly Ser Ala Val Val Ser Leu Val Ile
 355 360 365
 Lys Trp Glu Tyr Val Leu Leu Leu Phe Leu Leu Leu Ala Asp Ala Arg
 370 375 380
 Ile Cys Ala Cys Leu Trp Met Met Leu Leu Ile Ala Gln Ala Glu Ala
 385 390 395 400
 Ala Leu Glu Asn Leu Val Val Leu Asn Ala Ala Val Ala Gly Ala
 405 410 415
 His Gly Thr Leu Ser Phe Leu Val Phe Phe Cys Ala Ala Trp Tyr Ile
 420 425 430
 Lys Gly Arg Leu Val Pro Gly Ala Ala Tyr Ala Phe Tyr Gly Val Trp
 435 440 445
 Pro Leu Leu Leu Leu Leu Leu Ala Leu Pro Pro Arg Ala Tyr Ala
 450 455 460

<210> 47

<211> 2082

<212> DNA

<213> Hepatitis C virus

<220>

<221> CDS

<222> 1..2079

<220>

<221> mat_peptide

<222> 1..2076

<400> 47

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gtg ggg tac att ccg ctc gtc ggc gcc ccc cta ggg ggc gct gcc agg				96
Val Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu Gly Gly Ala Ala Arg	20	25	30	
gcc ctg gcg cat ggc gtc cgg gtt ctg gag gac ggc gtg aac tat gca				144
Ala Leu Ala His Gly Val Arg Val Leu Glu Asp Gly Val Asn Tyr Ala	35	40	45	
aca ggg aat ttg ccc ggt tgc tct ttc tct atc ttc ctc ttg gct ttg				192
Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu	50	55	60	
ctg tcc tgt ctg acc gtt cca gct tcc gct tat gaa gtg cgc aac gtg				240
Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Tyr Glu Val Arg Asn Val	65	70	75	80
tcc ggg atg tac cat gtc acg aac gac tgc tcc aac tca agc att gtg				288
Ser Gly Met Tyr His Val Thr Asn Asp Cys Ser Asn Ser Ser Ile Val	85	90	95	
tat gag gca gcg gac atg atc atg cac acc ccc ggg tgc gtg ccc tgc				336
Tyr Glu Ala Ala Asp Met Ile Met His Thr Pro Gly Cys Val Pro Cys	100	105	110	
gtt cgg gag aac aac tct tcc cgc tgc tgg gta gcg ctc acc ccc acg				384
Val Arg Glu Asn Asn Ser Ser Arg Cys Trp Val Ala Leu Thr Pro Thr	115	120	125	
ctc gca gct agg aac gcc agc gtc ccc acc acg aca ata cga cgc cac				432
Leu Ala Ala Arg Asn Ala Ser Val Pro Thr Thr Thr Ile Arg Arg His	130	135	140	
gtc gat ttg ctc gtt ggg gcg gct gct ttc tgt tcc gct atg tac gtg				480
Val Asp Leu Leu Val Gly Ala Ala Phe Cys Ser Ala Met Tyr Val	145	150	155	160
ggg gac ctc tgc gga tct gtc ttc ctc gtc tcc cag ctg ttc acc atc				528
Gly Asp Leu Cys Gly Ser Val Phe Leu Val Ser Gln Leu Phe Thr Ile	165	170	175	
tcg cct cgc cgg cat gag acg gtg cag gac tgc aat tgc tca atc tat				576
Ser Pro Arg Arg His Glu Thr Val Gln Asp Cys Asn Cys Ser Ile Tyr	180	185	190	
ccc ggc cac ata acg ggt cac cgt atg gct tgg gat atg atg atg aac				624
Pro Gly His Ile Thr Gly His Arg Met Ala Trp Asp Met Met Met Asn	195	200	205	
tgg tcg cct aca acg gcc ctg gtg gta tcg cag ctg ctc cgg atc cca				672
Trp Ser Pro Thr Thr Ala Leu Val Val Ser Gln Leu Leu Arg Ile Pro	210	215	220	
caa gct gtc gtg gac atg gtg gcg ggg gcc cat tgg gga gtc ctg gcg				720
Gln Ala Val Val Asp Met Val Ala Gly Ala His Trp Gly Val Leu Ala	225	230	235	240
ggc ctc gcc tac tat tcc atg gtg ggg aac tgg gct aag gtt ttg gtt				768
Gly Leu Ala Tyr Tyr Ser Met Val Gly Asn Trp Ala Lys Val Leu Val	245	250	255	
gtg atg cta ctc ttt gcc ggc gtc gac ggg cat acc cgc gtg tca gga				816

Val	Met	Leu	Leu	Phe	Ala	Gly	Val	Asp	Gly	His	Thr	Arg	Val	Ser	Gly		
			260					265					270				
ggg	gca	gca	gcc	tcc	gat	acc	agg	ggc	ctt	gtg	tcc	ctc	ttt	agc	ccc	864	
Gly	Ala	Ala	Ala	Ser	Asp	Thr	Arg	Gly	Leu	Val	Ser	Leu	Phe	Ser	Pro		
		275					280					285					
ggg	tcg	gct	cag	aaa	atc	cag	ctc	gta	aac	acc	aac	ggc	agt	tgg	cac	912	
Gly	Ser	Ala	Gln	Lys	Ile	Gln	Leu	Val	Asn	Thr	Asn	Gly	Ser	Trp	His		
	290					295					300						
atc	aac	agg	act	gcc	ctg	aac	tgc	aac	gac	tcc	ctc	caa	aca	ggg	ttc	960	
Ile	Asn	Arg	Thr	Ala		Leu	Asn	Cys	Asn	Asp	Ser	Leu	Gln	Thr	Gly	Phe	
305					310					315					320		
ttt	gcc	gca	cta	ttc	tac	aaa	cac	aaa	ttc	aac	tcg	tct	gga	tgc	cca	1008	
Phe	Ala	Ala	Leu	Phe	Tyr	Lys	His	Lys	Phe	Asn	Ser	Ser	Gly	Cys	Pro		
				325					330					335			
gag	cgc	ttg	gcc	agc	tgt	cgc	tcc	atc	gac	aag	ttc	gct	cag	ggg	tgg	1056	
Glu	Arg	Leu	Ala	Ser	Cys	Arg	Ser	Ile	Asp	Lys	Phe	Ala	Gln	Gly	Trp		
		340					345						350				
ggt	ccc	ctc	act	tac	act	gag	cct	aac	agc	tcg	gac	cag	agg	ccc	tac	1104	
Gly	Pro	Leu	Thr	Tyr	Thr	Glu	Pro	Asn	Ser	Ser	Asp	Gln	Arg	Pro	Tyr		
		355				360						365					
tgc	tgg	cac	tac	gcg	cct	cga	ccg	tgt	ggt	att	gta	ccc	gcg	tct	cag	1152	
Cys	Trp	His	Tyr	Ala	Pro	Arg	Pro	Cys	Gly	Ile	Val	Pro	Ala	Ser	Gln		
	370					375					380						
gtg	tgc	ggt	cca	gtg	tat	tgc	ttc	acc	ccg	agc	cct	gtt	gtg	gtg	ggg	1200	
Val	Cys	Gly	Pro	Val	Tyr	Cys	Phe	Thr	Pro	Ser	Pro	Val	Val	Val	Gly		
385					390					395					400		
acg	acc	gat	cgg	ttt	ggt	gtc	ccc	acg	tat	aac	tgg	ggg	gcg	aac	gac	1248	
Thr	Thr	Asp	Arg	Phe	Gly	Val	Pro	Thr	Tyr	Asn	Trp	Gly	Ala	Asn	Asp		
				405					410					415			
tcg	gat	gtg	ctg	att	ctc	aac	aac	acg	cgg	ccg	ccg	cga	ggc	aac	tgg	1296	
Ser	Asp	Val	Leu	Ile	Leu	Asn	Asn	Thr	Arg	Pro	Pro	Arg	Gly	Asn	Trp		
		420						425					430				
ttc	ggc	tgt	aca	tgg	atg	aat	ggc	act	ggg	ttc	acc	aag	acg	tgt	ggg	1344	
Phe	Gly	Cys	Thr	Trp	Met	Asn	Gly	Thr	Gly	Phe	Thr	Lys	Thr	Cys	Gly		
		435					440					445					
ggc	ccc	ccg	tgc	aac	atc	ggg	ggg	gcc	ggc	aac	aac	acc	ttg	acc	tgc	1392	
Gly	Pro	Pro	Cys	Asn	Ile	Gly	Gly	Ala	Gly	Asn	Asn	Thr	Leu	Thr	Cys		
	450					455					460						
ccc	act	gac	tgt	ttt	cgg	aag	cac	ccc	gag	gcc	acc	tac	gcc	aga	tgc	1440	
Pro	Thr	Asp	Cys	Phe	Arg	Lys	His	Pro	Glu	Ala	Thr	Tyr	Ala	Arg	Cys		
465					470					475					480		
ggt	tct	ggg	ccc	tgg	ctg	aca	cct	agg	tgt	atg	gtt	cat	tac	cca	tat	1488	
Gly	Ser	Gly	Pro	Trp	Leu	Thr	Pro	Arg	Cys	Met	Val	His	Tyr	Pro	Tyr		
				485					490					495			
agg	ctc	tgg	cac	tac	ccc	tgc	act	gtc	aac	ttc	acc	atc	ttc	aag	gtt	1536	
Arg	Leu	Trp	His	Tyr	Pro	Cys	Thr	Val	Asn	Phe	Thr	Ile	Phe	Lys	Val		
			500					505					510				
agg	atg	tac	gtg	ggg	ggc	gtg	gag	cac	agg	ttc	gaa	gcc	gca	tgc	aat	1584	

Arg	Met	Tyr	Val	Gly	Gly	Val	Glu	His	Arg	Phe	Glu	Ala	Ala	Cys	Asn		
		515					520					525					
tgg	act	cga	gga	gag	cgt	tgt	gac	ttg	gag	gac	agg	gat	aga	tca	gag	1632	
Trp	Thr	Arg	Gly	Glu	Arg	Cys	Asp	Leu	Glu	Asp	Arg	Asp	Arg	Ser	Glu		
	530					535				540							
ctt	agc	ccg	ctg	ctg	ctg	tct	aca	aca	gag	tgg	cag	ata	ctg	ccc	tgt	1680	
Leu	Ser	Pro	Leu	Leu	Leu	Ser	Thr	Thr	Glu	Trp	Gln	Ile	Leu	Pro	Cys		
	545				550				555						560		
tcc	ttc	acc	acc	ctg	ccg	gcc	cta	tcc	acc	ggc	ctg	atc	cac	ctc	cat	1728	
Ser	Phe	Thr	Thr		Leu	Pro	Ala	Leu	Ser	Thr	Gly	Leu	Ile	His	Leu	His	
				565				570						575			
cag	aac	atc	gtg	gac	gtg	caa	tac	ctg	tac	ggc	gta	ggg	tcg	gcg	gtt	1776	
Gln	Asn	Ile	Val	Asp	Val	Gln	Tyr	Leu	Tyr	Gly	Val	Gly	Ser	Ala	Val		
			580					585					590				
gtc	tcc	ctt	gtc	atc	aaa	tgg	gag	tat	gtc	ctg	ttg	ctc	ttc	ctt	ctc	1824	
Val	Ser	Leu	Val	Ile	Lys	Trp	Glu	Tyr	Val	Leu	Leu	Leu	Phe	Leu	Leu		
		595				600					605						
ctg	gca	gac	gcg	cgc	atc	tgc	gcc	tgc	tta	tgg	atg	atg	ctg	ctg	ata	1872	
Leu	Ala	Asp	Ala	Arg	Ile	Cys	Ala	Cys	Leu	Trp	Met	Met	Leu	Leu	Ile		
	610					615				620							
gct	caa	gct	gag	gcc	gcc	tta	gag	aac	ctg	gtg	gtc	ctc	aat	gcg	gcg	1920	
Ala	Gln	Ala	Glu	Ala	Ala	Leu	Glu	Asn	Leu	Val	Val	Leu	Asn	Ala	Ala		
	625				630				635					640			
gcc	gtg	gcc	ggg	gcg	cat	ggc	act	ctt	tcc	ttc	ctt	gtg	ttc	ttc	tgt	1968	
Ala	Val	Ala	Gly	Ala	His	Gly	Thr	Leu	Ser	Phe	Leu	Val	Phe	Phe	Cys		
			645					650					655				
gct	gcc	tgg	tac	atc	aag	ggc	agg	ctg	gtc	cct	ggc	gcg	gca	tac	gcc	2016	
Ala	Ala	Trp	Tyr	Ile	Lys	Gly	Arg	Leu	Val	Pro	Gly	Ala	Ala	Tyr	Ala		
		660						665				670					
ttc	tat	ggc	gtg	tgg	ccg	ctg	ctc	ctg	ctt	ctg	ctg	gcc	tta	cca	cca	2064	
Phe	Tyr	Gly	Val	Trp	Pro	Leu	Leu	Leu	Leu	Leu	Ala	Leu	Pro	Pro			
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cga	gct	tat	gcc	tagtaa												2082	
Arg	Ala	Tyr	Ala														
	690																

<210> 48

<211> 692

<212> PRT

<213> Hepatitis C virus

<400> 48

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		20					25					30					
Ala	Leu	Ala	His	Gly	Val	Arg	Val	Leu	Glu	Asp	Gly	Val	Asn	Tyr	Ala		

35					40					45					
Thr	Gly	Asn	Leu	Pro	Gly	Cys	Ser	Phe	Ser	Ile	Phe	Leu	Leu	Ala	Leu
50						55					60				
Leu	Ser	Cys	Leu	Thr	Val	Pro	Ala	Ser	Ala	Tyr	Glu	Val	Arg	Asn	Val
65					70					75					80
Ser	Gly	Met	Tyr	His	Val	Thr	Asn	Asp	Cys	Ser	Asn	Ser	Ser	Ile	Val
				85					90					95	
Tyr	Glu	Ala	Ala	Asp	Met	Ile	Met	His	Thr	Pro	Gly	Cys	Val	Pro	Cys
			100					105					110		
Val	Arg	Glu	Asn	Asn	Ser	Ser	Arg	Cys	Trp	Val	Ala	Leu	Thr	Pro	Thr
			115				120					125			
Leu	Ala	Ala	Arg	Asn	Ala	Ser	Val	Pro	Thr	Thr	Thr	Ile	Arg	Arg	His
130						135						140			
Val	Asp	Leu	Leu	Val	Gly	Ala	Ala	Ala	Phe	Cys	Ser	Ala	Met	Tyr	Val
145					150					155					160
Gly	Asp	Leu	Cys	Gly	Ser	Val	Phe	Leu	Val	Ser	Gln	Leu	Phe	Thr	Ile
				165					170					175	
Ser	Pro	Arg	Arg	His	Glu	Thr	Val	Gln	Asp	Cys	Asn	Cys	Ser	Ile	Tyr
			180					185					190		
Pro	Gly	His	Ile	Thr	Gly	His	Arg	Met	Ala	Trp	Asp	Met	Met	Met	Asn
		195					200					205			
Trp	Ser	Pro	Thr	Thr	Ala	Leu	Val	Val	Ser	Gln	Leu	Leu	Arg	Ile	Pro
210						215					220				
Gln	Ala	Val	Val	Asp	Met	Val	Ala	Gly	Ala	His	Trp	Gly	Val	Leu	Ala
225					230					235					240
Gly	Leu	Ala	Tyr	Tyr	Ser	Met	Val	Gly	Asn	Trp	Ala	Lys	Val	Leu	Val
				245					250					255	
Val	Met	Leu	Leu	Phe	Ala	Gly	Val	Asp	Gly	His	Thr	Arg	Val	Ser	Gly
			260					265					270		
Gly	Ala	Ala	Ala	Ser	Asp	Thr	Arg	Gly	Leu	Val	Ser	Leu	Phe	Ser	Pro
			275				280						285		
Gly	Ser	Ala	Gln	Lys	Ile	Gln	Leu	Val	Asn	Thr	Asn	Gly	Ser	Trp	His
290						295					300				
Ile	Asn	Arg	Thr	Ala	Leu	Asn	Cys	Asn	Asp	Ser	Leu	Gln	Thr	Gly	Phe
305						310					315				320
Phe	Ala	Ala	Leu	Phe	Tyr	Lys	His	Lys	Phe	Asn	Ser	Ser	Gly	Cys	Pro
				325					330					335	
Glu	Arg	Leu	Ala	Ser	Cys	Arg	Ser	Ile	Asp	Lys	Phe	Ala	Gln	Gly	Trp
			340					345					350		
Gly	Pro	Leu	Thr	Tyr	Thr	Glu	Pro	Asn	Ser	Ser	Asp	Gln	Arg	Pro	Tyr
			355				360					365			
Cys	Trp	His	Tyr	Ala	Pro	Arg	Pro	Cys	Gly	Ile	Val	Pro	Ala	Ser	Gln
370						375					380				

Val Cys Gly Pro Val Tyr Cys Phe Thr Pro Ser Pro Val Val Val Gly
 385 390 395 400
 Thr Thr Asp Arg Phe Gly Val Pro Thr Tyr Asn Trp Gly Ala Asn Asp
 405 410 415
 Ser Asp Val Leu Ile Leu Asn Asn Thr Arg Pro Pro Arg Gly Asn Trp
 420 425 430
 Phe Gly Cys Thr Trp Met Asn Gly Thr Gly Phe Thr Lys Thr Cys Gly
 435 440 445
 Gly Pro Pro Cys Asn Ile Gly Gly Ala Gly Asn Asn Thr Leu Thr Cys
 450 455 460
 Pro Thr Asp Cys Phe Arg Lys His Pro Glu Ala Thr Tyr Ala Arg Cys
 465 470 475 480
 Gly Ser Gly Pro Trp Leu Thr Pro Arg Cys Met Val His Tyr Pro Tyr
 485 490 495
 Arg Leu Trp His Tyr Pro Cys Thr Val Asn Phe Thr Ile Phe Lys Val
 500 505 510
 Arg Met Tyr Val Gly Gly Val Glu His Arg Phe Glu Ala Ala Cys Asn
 515 520 525
 Trp Thr Arg Gly Glu Arg Cys Asp Leu Glu Asp Arg Asp Arg Ser Glu
 530 535 540
 Leu Ser Pro Leu Leu Leu Ser Thr Thr Glu Trp Gln Ile Leu Pro Cys
 545 550 555 560
 Ser Phe Thr Thr Leu Pro Ala Leu Ser Thr Gly Leu Ile His Leu His
 565 570 575
 Gln Asn Ile Val Asp Val Gln Tyr Leu Tyr Gly Val Gly Ser Ala Val
 580 585 590
 Val Ser Leu Val Ile Lys Trp Glu Tyr Val Leu Leu Leu Phe Leu Leu
 595 600 605
 Leu Ala Asp Ala Arg Ile Cys Ala Cys Leu Trp Met Met Leu Leu Ile
 610 615 620
 Ala Gln Ala Glu Ala Ala Leu Glu Asn Leu Val Val Leu Asn Ala Ala
 625 630 635 640
 Ala Val Ala Gly Ala His Gly Thr Leu Ser Phe Leu Val Phe Phe Cys
 645 650 655
 Ala Ala Trp Tyr Ile Lys Gly Arg Leu Val Pro Gly Ala Ala Tyr Ala
 660 665 670
 Phe Tyr Gly Val Trp Pro Leu Leu Leu Leu Leu Ala Leu Pro Pro
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 Arg Ala Tyr Ala
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<211> 2433
 <212> DNA
 <213> Hepatitis C virus

<220>
 <221> CDS
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<220>
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<400> 49

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cgc cgc cca cag gac gtc aag ttc ccg ggc ggt ggt cag atc gtt ggt	96
Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly	
20 25 30	
gga gtt tac ctg ttg ccg cgc agg ggc ccc agg ttg ggt gtg cgc gcg	144
Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala	
35 40 45	
act agg aag act tcc gag cgg tcg caa cct cgt ggg agg cga caa cct	192
Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro	
50 55 60	
atc ccc aag gct cgc cga ccc gag ggt agg gcc tgg gct cag ccc ggg	240
Ile Pro Lys Ala Arg Arg Pro Glu Gly Arg Ala Trp Ala Gln Pro Gly	
65 70 75 80	
tac cct tgg ccc ctc tat ggc aat gag ggc atg ggg tgg gca gga tgg	288
Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Met Gly Trp Ala Gly Trp	
85 90 95	
ctc ctg tca ccc cgc ggc tct cgg cct agt tgg ggc cct aca gac ccc	336
Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro	
100 105 110	
cgg cgt agg tcg cgt aat ttg ggt aag gtc atc gat acc ctt aca tgc	384
Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys	
115 120 125	
ggc ttc gcc gac ctc gtg ggg tac att ccg ctc gtc ggc gcc ccc cta	432
Gly Phe Ala Asp Leu Val Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu	
130 135 140	
ggg ggc gct gcc agg gcc ctg gcg cat ggc gtc cgg gtt ctg gag gac	480
Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp	
145 150 155 160	
ggc gtg aac tat gca aca ggg aat ttg ccc ggt tgc tct ttc tct atc	528
Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile	
165 170 175	
ttc ctc ttg gct ttg ctg tcc tgt ctg acc gtt cca gct tcc gct tat	576
Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Tyr	
180 185 190	
gaa gtg cgc aac gtg tcc ggg atg tac cat gtc acg aac gac tgc tcc	624

Glu	Val	Arg	Asn	Val	Ser	Gly	Met	Tyr	His	Val	Thr	Asn	Asp	Cys	Ser		
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aac	tca	agc	att	gtg	tat	gag	gca	gcg	gac	atg	atc	atg	cac	acc	ccc	672	
Asn	Ser	Ser	Ile	Val	Tyr	Glu	Ala	Ala	Asp	Met	Ile	Met	His	Thr	Pro		
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ggg	tgc	gtg	ccc	tgc	gtt	cgg	gag	aac	aac	tct	tcc	cgc	tgc	tgg	gta	720	
Gly	Cys	Val	Pro	Cys	Val	Arg	Glu	Asn	Asn	Ser	Ser	Arg	Cys	Trp	Val		
	225				230					235					240		
gcg	ctc	acc	ccc	acg	ctc	gca	gct	agg	aac	gcc	agc	gtc	ccc	acc	acg	768	
Ala	Leu	Thr	Pro	Thr	Leu	Ala	Ala	Arg	Asn	Ala	Ser	Val	Pro	Thr	Thr		
				245					250					255			
aca	ata	cga	cgc	cac	gtc	gat	ttg	ctc	gtt	ggg	gcg	gct	gct	ttc	tgt	816	
Thr	Ile	Arg	Arg	His	Val	Asp	Leu	Leu	Val	Gly	Ala	Ala	Ala	Phe	Cys		
			260					265					270				
tcc	gct	atg	tac	gtg	ggg	gac	ctc	tgc	gga	tct	gtc	ttc	ctc	gtc	tcc	864	
Ser	Ala	Met	Tyr	Val	Gly	Asp	Leu	Cys	Gly	Ser	Val	Phe	Leu	Val	Ser		
	275					280						285					
cag	ctg	ttc	acc	atc	tcg	cct	cgc	cgg	cat	gag	acg	gtg	cag	gac	tgc	912	
Gln	Leu	Phe	Thr	Ile	Ser	Pro	Arg	Arg	His	Glu	Thr	Val	Gln	Asp	Cys		
	290					295					300						
aat	tgc	tca	atc	tat	ccc	ggc	cac	ata	acg	ggc	cac	cgt	atg	gct	tgg	960	
Asn	Cys	Ser	Ile	Tyr	Pro	Gly	His	Ile	Thr	Gly	His	Arg	Met	Ala	Trp		
	305				310					315					320		
gat	atg	atg	atg	aac	tgg	tcg	cct	aca	acg	gcc	ctg	gtg	gta	tcg	cag	1008	
Asp	Met	Met	Met	Asn	Trp	Ser	Pro	Thr	Thr	Ala	Leu	Val	Val	Ser	Gln		
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ctg	ctc	cgg	atc	cca	caa	gct	gtc	gtg	gac	atg	gtg	gcg	ggg	gcc	cat	1056	
Leu	Leu	Arg	Ile	Pro	Gln	Ala	Val	Val	Asp	Met	Val	Ala	Gly	Ala	His		
			340					345					350				
tgg	gga	gtc	ctg	gcg	ggc	ctc	gcc	tac	tat	tcc	atg	gtg	ggg	aac	tgg	1104	
Trp	Gly	Val	Leu	Ala	Gly	Leu	Ala	Tyr	Tyr	Ser	Met	Val	Gly	Asn	Trp		
	355					360						365					
gct	aag	gtt	ttg	gtt	gtg	atg	cta	ctc	ttt	gcc	ggc	gtc	gac	ggg	cat	1152	
Ala	Lys	Val	Leu	Val	Val	Met	Leu	Leu	Phe	Ala	Gly	Val	Asp	Gly	His		
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acc	cgc	gtg	tca	gga	ggg	gca	gca	gcc	tcc	gat	acc	agg	ggc	ctt	gtg	1200	
Thr	Arg	Val	Ser	Gly	Gly	Ala	Ala	Ala	Ser	Asp	Thr	Arg	Gly	Leu	Val		
	385				390					395					400		
tcc	ctc	ttt	agc	ccc	ggg	tcg	gct	cag	aaa	atc	cag	ctc	gta	aac	acc	1248	
Ser	Leu	Phe	Ser	Pro	Gly	Ser	Ala	Gln	Lys	Ile	Gln	Leu	Val	Asn	Thr		
				405				410						415			
aac	ggc	agt	tgg	cac	atc	aac	agg	act	gcc	ctg	aac	tgc	aac	gac	tcc	1296	
Asn	Gly	Ser	Trp	His	Ile	Asn	Arg	Thr	Ala	Leu	Asn	Cys	Asn	Asp	Ser		
			420					425					430				
ctc	caa	aca	ggg	ttc	ttt	gcc	gca	cta	ttc	tac	aaa	cac	aaa	ttc	aac	1344	
Leu	Gln	Thr	Gly	Phe	Phe	Ala	Ala	Leu	Phe	Tyr	Lys	His	Lys	Phe	Asn		
		435				440						445					

tcg tct gga tgc cca gag cgc ttg gcc agc tgt cgc tcc atc gac aag Ser Ser Gly Cys Pro Glu Arg Leu Ala Ser Cys Arg Ser Ile Asp Lys 450 455 460	1392
ttc gct cag ggg tgg ggt ccc ctc act tac act gag cct aac agc tcg Phe Ala Gln Gly Trp Gly Pro Leu Thr Tyr Thr Glu Pro Asn Ser Ser 465 470 475 480	1440
gac cag agg ccc tac tgc tgg cac tac gcg cct cga ccg tgt ggt att Asp Gln Arg Pro Tyr Cys Trp His Tyr Ala Pro Arg Pro Cys Gly Ile 485 490 495	1488
gta ccc gcg tct cag gtg tgc ggt cca gtg tat tgc ttc acc ccg agc Val Pro Ala Ser Gln Val Cys Gly Pro Val Tyr Cys Phe Thr Pro Ser 500 505 510	1536
cct gtt gtg gtg ggg acg acc gat cgg ttt ggt gtc ccc acg tat aac Pro Val Val Val Gly Thr Thr Asp Arg Phe Gly Val Pro Thr Tyr Asn 515 520 525	1584
tgg ggg gcg aac gac tcg gat gtg ctg att ctc aac aac acg cgg ccg Trp Gly Ala Asn Asp Ser Asp Val Leu Ile Leu Asn Asn Thr Arg Pro 530 535 540	1632
ccg cga ggc aac tgg ttc ggc tgt aca tgg atg aat ggc act ggg ttc Pro Arg Gly Asn Trp Phe Gly Cys Thr Trp Met Asn Gly Thr Gly Phe 545 550 555 560	1680
acc aag acg tgt ggg ggc ccc ccg tgc aac atc ggg ggg gcc ggc aac Thr Lys Thr Cys Gly Gly Pro Pro Cys Asn Ile Gly Gly Ala Gly Asn 565 570 575	1728
aac acc ttg acc tgc ccc act gac tgt ttt cgg aag cac ccc gag gcc Asn Thr Leu Thr Cys Pro Thr Asp Cys Phe Arg Lys His Pro Glu Ala 580 585 590	1776
acc tac gcc aga tgc ggt tct ggg ccc tgg ctg aca cct agg tgt atg Thr Tyr Ala Arg Cys Gly Ser Gly Pro Trp Leu Thr Pro Arg Cys Met 595 600 605	1824
gtt cat tac cca tat agg ctc tgg cac tac ccc tgc act gtc aac ttc Val His Tyr Pro Tyr Arg Leu Trp His Tyr Pro Cys Thr Val Asn Phe 610 615 620	1872
acc atc ttc aag gtt agg atg tac gtg ggg ggc gtg gag cac agg ttc Thr Ile Phe Lys Val Arg Met Tyr Val Gly Gly Val Glu His Arg Phe 625 630 635 640	1920
gaa gcc gca tgc aat tgg act cga gga gag cgt tgt gac ttg gag gac Glu Ala Ala Cys Asn Trp Thr Arg Gly Glu Arg Cys Asp Leu Glu Asp 645 650 655	1968
agg gat aga tca gag ctt agc ccg ctg ctg ctg tct aca aca gag tgg Arg Asp Arg Ser Glu Leu Ser Pro Leu Leu Leu Ser Thr Thr Glu Trp 660 665 670	2016
cag ata ctg ccc tgt tcc ttc acc acc ctg ccg gcc cta tcc acc ggc Gln Ile Leu Pro Cys Ser Phe Thr Thr Leu Pro Ala Leu Ser Thr Gly 675 680 685	2064
ctg atc cac ctc cat cag aac atc gtg gac gtg caa tac ctg tac ggt Leu Ile His Leu His Gln Asn Ile Val Asp Val Gln Tyr Leu Tyr Gly 690 695 700	2112

gta ggg tgc gcg gtt gtc tcc ctt gtc atc aaa tgg gag tat gtc ctg 2160
 Val Gly Ser Ala Val Val Ser Leu Val Ile Lys Trp Glu Tyr Val Leu
 705 710 715 720

ttg ctc ttc ctt ctc ctg gca gac gcg cgc atc tgc gcc tgc tta tgg 2208
 Leu Leu Phe Leu Leu Leu Ala Asp Ala Arg Ile Cys Ala Cys Leu Trp
 725 730 735

atg atg ctg ctg ata gct caa gct gag gcc gcc tta gag aac ctg gtg 2256
 Met Met Leu Leu Ile Ala Gln Ala Glu Ala Ala Leu Glu Asn Leu Val
 740 745 750

gtc ctc aat gcg gcg gcc gtg gcc ggg gcg cat ggc act ctt tcc ttc 2304
 Val Leu Asn Ala Ala Ala Val Ala Gly Ala His Gly Thr Leu Ser Phe
 755 760 765

ctt gtg ttc ttc tgt gct gcc tgg tac atc aag ggc agg ctg gtc cct 2352
 Leu Val Phe Phe Cys Ala Ala Trp Tyr Ile Lys Gly Arg Leu Val Pro
 770 775 780

ggc gcg gca tac gcc ttc tat ggc gtg tgg ccg ctg ctc ctg ctt ctg 2400
 Gly Ala Ala Tyr Ala Phe Tyr Gly Val Trp Pro Leu Leu Leu Leu Leu
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<210> 50

<211> 809

<212> PRT

<213> Hepatitis C virus

<400> 50

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn
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Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly
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Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala
 35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro
 50 55 60

Ile Pro Lys Ala Arg Arg Pro Glu Gly Arg Ala Trp Ala Gln Pro Gly
 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Met Gly Trp Ala Gly Trp
 85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
 100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
 115 120 125

Gly Phe Ala Asp Leu Val Gly Tyr Ile Pro Leu Val Gly Ala Pro Leu
 130 135 140

Gly Gly Ala Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp
 145 150 155 160
 Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile
 165 170 175
 Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Tyr
 180 185 190
 Glu Val Arg Asn Val Ser Gly Met Tyr His Val Thr Asn Asp Cys Ser
 195 200 205
 Asn Ser Ser Ile Val Tyr Glu Ala Ala Asp Met Ile Met His Thr Pro
 210 215 220
 Gly Cys Val Pro Cys Val Arg Glu Asn Asn Ser Ser Arg Cys Trp Val
 225 230 235 240
 Ala Leu Thr Pro Thr Leu Ala Ala Arg Asn Ala Ser Val Pro Thr Thr
 245 250 255
 Thr Ile Arg Arg His Val Asp Leu Leu Val Gly Ala Ala Ala Phe Cys
 260 265 270
 Ser Ala Met Tyr Val Gly Asp Leu Cys Gly Ser Val Phe Leu Val Ser
 275 280 285
 Gln Leu Phe Thr Ile Ser Pro Arg Arg His Glu Thr Val Gln Asp Cys
 290 295 300
 Asn Cys Ser Ile Tyr Pro Gly His Ile Thr Gly His Arg Met Ala Trp
 305 310 315 320
 Asp Met Met Met Asn Trp Ser Pro Thr Thr Ala Leu Val Val Ser Gln
 325 330 335
 Leu Leu Arg Ile Pro Gln Ala Val Val Asp Met Val Ala Gly Ala His
 340 345 350
 Trp Gly Val Leu Ala Gly Leu Ala Tyr Tyr Ser Met Val Gly Asn Trp
 355 360 365
 Ala Lys Val Leu Val Val Met Leu Leu Phe Ala Gly Val Asp Gly His
 370 375 380
 Thr Arg Val Ser Gly Gly Ala Ala Ala Ser Asp Thr Arg Gly Leu Val
 385 390 395 400
 Ser Leu Phe Ser Pro Gly Ser Ala Gln Lys Ile Gln Leu Val Asn Thr
 405 410 415
 Asn Gly Ser Trp His Ile Asn Arg Thr Ala Leu Asn Cys Asn Asp Ser
 420 425 430
 Leu Gln Thr Gly Phe Phe Ala Ala Leu Phe Tyr Lys His Lys Phe Asn
 435 440 445
 Ser Ser Gly Cys Pro Glu Arg Leu Ala Ser Cys Arg Ser Ile Asp Lys
 450 455 460
 Phe Ala Gln Gly Trp Gly Pro Leu Thr Tyr Thr Glu Pro Asn Ser Ser
 465 470 475 480

Asp Gln Arg Pro Tyr Cys Trp His Tyr Ala Pro Arg Pro Cys Gly Ile
 485 490 495
 Val Pro Ala Ser Gln Val Cys Gly Pro Val Tyr Cys Phe Thr Pro Ser
 500 505 510
 Pro Val Val Val Gly Thr Thr Asp Arg Phe Gly Val Pro Thr Tyr Asn
 515 520 525
 Trp Gly Ala Asn Asp Ser Asp Val Leu Ile Leu Asn Asn Thr Arg Pro
 530 535 540
 Pro Arg Gly Asn Trp Phe Gly Cys Thr Trp Met Asn Gly Thr Gly Phe
 545 550 555 560
 Thr Lys Thr Cys Gly Gly Pro Pro Cys Asn Ile Gly Gly Ala Gly Asn
 565 570 575
 Asn Thr Leu Thr Cys Pro Thr Asp Cys Phe Arg Lys His Pro Glu Ala
 580 585 590
 Thr Tyr Ala Arg Cys Gly Ser Gly Pro Trp Leu Thr Pro Arg Cys Met
 595 600 605
 Val His Tyr Pro Tyr Arg Leu Trp His Tyr Pro Cys Thr Val Asn Phe
 610 615 620
 Thr Ile Phe Lys Val Arg Met Tyr Val Gly Gly Val Glu His Arg Phe
 625 630 635 640
 Glu Ala Ala Cys Asn Trp Thr Arg Gly Glu Arg Cys Asp Leu Glu Asp
 645 650 655
 Arg Asp Arg Ser Glu Leu Ser Pro Leu Leu Leu Ser Thr Thr Glu Trp
 660 665 670
 Gln Ile Leu Pro Cys Ser Phe Thr Thr Leu Pro Ala Leu Ser Thr Gly
 675 680 685
 Leu Ile His Leu His Gln Asn Ile Val Asp Val Gln Tyr Leu Tyr Gly
 690 695 700
 Val Gly Ser Ala Val Val Ser Leu Val Ile Lys Trp Glu Tyr Val Leu
 705 710 715 720
 Leu Leu Phe Leu Leu Leu Ala Asp Ala Arg Ile Cys Ala Cys Leu Trp
 725 730 735
 Met Met Leu Leu Ile Ala Gln Ala Glu Ala Ala Leu Glu Asn Leu Val
 740 745 750
 Val Leu Asn Ala Ala Ala Val Ala Gly Ala His Gly Thr Leu Ser Phe
 755 760 765
 Leu Val Phe Phe Cys Ala Ala Trp Tyr Ile Lys Gly Arg Leu Val Pro
 770 775 780
 Gly Ala Ala Tyr Ala Phe Tyr Gly Val Trp Pro Leu Leu Leu Leu Leu
 785 790 795 800
 Leu Ala Leu Pro Pro Arg Ala Tyr Ala
 805

<210> 51

<211> 17

<212> PRT

<213> Hepatitis C virus

<400> 51

Ser	Asn	Ser	Ser	Glu	Ala	Ala	Asp	Met	Ile	Met	His	Thr	Pro	Gly	Cys
1				5					10					15	

Val

<210> 52

<211> 22

<212> PRT

<213> Hepatitis C virus

<400> 52

Gly	Gly	Ile	Thr	Gly	His	Arg	Met	Ala	Trp	Asp	Met	Met	Met	Asn	Trp
1				5					10					15	

Ser	Pro	Thr	Thr	Ala	Leu
					20

<210> 53

<211> 37

<212> PRT

<213> Hepatitis C virus

<400> 53

Tyr	Glu	Val	Arg	Asn	Val	Ser	Gly	Ile	Tyr	His	Val	Thr	Asn	Asp	Cys
1				5					10					15	

Ser	Asn	Ser	Ser	Ile	Val	Tyr	Glu	Ala	Ala	Asp	Met	Ile	Met	His	Thr
				20				25					30		

Pro	Gly	Cys	Gly	Lys
				35

<210> 54

<211> 25

<212> PRT

<213> Hepatitis C virus

<400> 54

Gly	Gly	Thr	Pro	Thr	Val	Ala	Thr	Arg	Asp	Gly	Lys	Leu	Pro	Ala	Thr
1				5					10					15	

Gln Leu Arg Arg His Ile Asp Leu Leu
 20 25

<210> 55

<211> 25

<212> PRT

<213> Hepatitis C virus

<400> 55

Gly Gly Thr Pro Thr Leu Ala Ala Arg Asp Ala Ser Val Pro Thr Thr
 1 5 10 15

Thr Ile Arg Arg His Val Asp Leu Leu
 20 25

<210> 56

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 56

Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Tyr Gln Val Arg Asn
 1 5 10 15

Ser Thr Gly Leu
 20

<210> 57

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 57

Gln Val Arg Asn Ser Thr Gly Leu Tyr His Val Thr Asn Asp Cys Pro
 1 5 10 15

Asn Ser Ser Ile
 20

<210> 58

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 58

Asn Asp Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala His Asp Ala Ile
 1 5 10 15

Leu His Thr Pro

20

<210> 59

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 59

Ser	Asn	Ser	Ser	Ile	Val	Tyr	Glu	Ala	Ala	Asp	Met	Ile	Met	His	Thr
1				5					10					15	

Pro	Gly	Cys	Val
			20

<210> 60

<211> 19

<212> PRT

<213> Hepatitis C virus

<400> 60

His	Asp	Ala	Ile	Leu	His	Thr	Pro	Gly	Val	Pro	Cys	Val	Arg	Glu	Gly
1				5					10					15	

Asn	Val	Ser
-----	-----	-----

<210> 61

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 61

Cys	Val	Arg	Glu	Gly	Asn	Val	Ser	Arg	Cys	Trp	Val	Ala	Met	Thr	Pro
1				5					10					15	

Thr	Val	Ala	Thr
			20

<210> 62

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 62

Ala	Met	Thr	Pro	Thr	Val	Ala	Thr	Arg	Asp	Gly	Lys	Leu	Pro	Ala	Thr
1				5					10					15	

Gln	Leu	Arg	Arg
-----	-----	-----	-----

20

<210> 63

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 63

Leu	Pro	Ala	Thr	Gln	Leu	Arg	Arg	His	Ile	Asp	Leu	Leu	Val	Gly	Ser
1				5					10					15	

Ala	Thr	Leu	Cys
			20

<210> 64

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 64

Leu	Val	Gly	Ser	Ala	Thr	Leu	Cys	Ser	Ala	Leu	Tyr	Val	Gly	Asp	Leu
1				5					10					15	

Cys	Gly	Ser	Val
			20

<210> 65

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 65

Gln	Leu	Phe	Thr	Phe	Ser	Pro	Arg	Arg	His	Trp	Thr	Thr	Gln	Gly	Cys
1				5					10					15	

Asn	Cys	Ser	Ile
			20

<210> 66

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 66

Thr	Gln	Gly	Cys	Asn	Cys	Ser	Ile	Tyr	Pro	Gly	His	Ile	Thr	Gly	His
1				5					10					15	

Arg	Met	Ala	Trp
-----	-----	-----	-----

20

<210> 67

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 67

Ile	Thr	Gly	His	Arg	Met	Ala	Trp	Asp	Met	Met	Met	Asn	Trp	Ser	Pro
1				5					10					15	

Thr	Ala	Ala	Leu
			20

<210> 68

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 68

Asn	Trp	Ser	Pro	Thr	Ala	Ala	Leu	Val	Met	Ala	Gln	Leu	Leu	Arg	Ile
1				5					10					15	

Pro	Gln	Ala	Ile
			20

<210> 69

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 69

Leu	Leu	Arg	Ile	Pro	Gln	Ala	Ile	Leu	Asp	Met	Ile	Ala	Gly	Ala	His
1				5					10					15	

Trp	Gly	Val	Leu
			20

<210> 70

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 70

Ala	Gly	Ala	His	Trp	Gly	Val	Leu	Ala	Gly	Ile	Ala	Tyr	Phe	Ser	Met
1				5					10					15	

Val	Gly	Asn	Met
-----	-----	-----	-----

20

<210> 71

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 71

Val	Val	Leu	Leu	Leu	Phe	Ala	Gly	Val	Asp	Ala	Glu	Thr	Ile	Val	Ser
1				5					10					15	

Gly	Gly	Gln	Ala
			20

<210> 72

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 72

Ser	Gly	Leu	Val	Ser	Leu	Phe	Thr	Pro	Gly	Ala	Lys	Gln	Asn	Ile	Gln
1				5					10					15	

Leu	Ile	Asn	Thr
			20

<210> 73

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 73

Gln	Asn	Ile	Gln	Leu	Ile	Asn	Thr	Asn	Gly	Gln	Trp	His	Ile	Asn	Ser
1				5					10					15	

Thr	Ala	Leu	Asn
			20

<210> 74

<211> 21

<212> PRT

<213> Hepatitis C virus

<400> 74

Leu	Asn	Cys	Asn	Glu	Ser	Leu	Asn	Thr	Gly	Trp	Trp	Leu	Ala	Gly	Leu
1				5					10					15	

Ile Tyr Gln His Lys
20

<210> 75

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 75

Ala Gly Leu Ile Tyr Gln His Lys Phe Asn Ser Ser Gly Cys Pro Glu
1 5 10 15

Arg Leu Ala Ser
20

<210> 76

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 76

Gly Cys Pro Glu Arg Leu Ala Ser Cys Arg Pro Leu Thr Asp Phe Asp
1 5 10 15

Gln Gly Trp Gly
20

<210> 77

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 77

Thr Asp Phe Asp Gln Gly Trp Gly Pro Ile Ser Tyr Ala Asn Gly Ser
1 5 10 15

Gly Pro Asp Gln
20

<210> 78

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 78

Ala Asn Gly Ser Gly Pro Asp Gln Arg Pro Tyr Cys Trp His Tyr Pro
1 5 10 15

Pro Lys Pro Cys
20

<210> 79

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 79

Trp His Tyr Pro Pro Lys Pro Cys Gly Ile Val Pro Ala Lys Ser Val
1 5 10 15

Cys Gly Pro Val
20

<210> 80

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 80

Ala Lys Ser Val Cys Gly Pro Val Tyr Cys Phe Thr Pro Ser Pro Val
1 5 10 15

Val Val Gly Thr
20

<210> 81

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 81

Pro Ser Pro Val Val Val Gly Thr Thr Asp Arg Ser Gly Ala Pro Thr
1 5 10 15

Tyr Ser Trp Gly
20

<210> 82

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 82

Gly Ala Pro Thr Tyr Ser Trp Gly Glu Asn Asp Thr Asp Val Phe Val
1 5 10 15

Leu Asn Asn Thr
20

<210> 83

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 83

Gly	Asn	Trp	Phe	Gly	Cys	Thr	Trp	Met	Asn	Ser	Thr	Gly	Phe	Thr	Lys
1				5					10					15	

Val Cys Gly Ala
20

<210> 84

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 84

Gly	Phe	Thr	Lys	Val	Cys	Gly	Ala	Pro	Pro	Val	Cys	Ile	Gly	Gly	Ala
1				5					10					15	

Gly Asn Asn Thr
20

<210> 85

<211> 19

<212> PRT

<213> Hepatitis C virus

<400> 85

Ile	Gly	Gly	Ala	Gly	Asn	Asn	Thr	Leu	His	Cys	Pro	Thr	Asp	Cys	Arg
1				5					10					15	

Lys His Pro

<210> 86

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 86

Thr	Asp	Cys	Phe	Arg	Lys	His	Pro	Asp	Ala	Thr	Tyr	Ser	Arg	Cys	Gly
1				5					10					15	

Ser Gly Pro Trp
20

<210> 87

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 87

Ser Arg Cys Gly Ser Gly Pro Trp Ile Thr Pro Arg Cys Leu Val Asp
1 5 10 15

Tyr Pro Tyr Arg
20

<210> 88

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 88

Cys Leu Val Asp Tyr Pro Tyr Arg Leu Trp His Tyr Pro Cys Thr Ile
1 5 10 15

Asn Tyr Thr Ile
20

<210> 89

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 89

Pro Cys Thr Ile Asn Tyr Thr Ile Phe Lys Ile Arg Met Tyr Val Gly
1 5 10 15

Gly Val Glu His
20

<210> 90

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 90

Met Tyr Val Gly Gly Val Glu His Arg Leu Glu Ala Ala Cys Asn Trp
1 5 10 15

Thr Pro Gly Glu
20

<210> 91

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 91

Ala Cys Asn Trp Thr Pro Gly Glu Arg Cys Asp Leu Glu Asp Arg Asp
1 5 10 15

Arg Ser Glu Leu
20

<210> 92

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 92

Glu Asp Arg Asp Arg Ser Glu Leu Ser Pro Leu Leu Leu Thr Thr Thr
1 5 10 15

Gln Trp Gln Val
20

<210> 93

<211> 9

<212> PRT

<213> Hepatitis C virus

<400> 93

Tyr Gln Val Arg Asn Ser Thr Gly Leu
1 5

<210> 94

<211> 29

<212> DNA

<213> Hepatitis C virus

<220>

<221> misc_feature

<223>/note="antisense"

<400> 94

acgtccgtac gttcgaatta attaatcga

<210> 95

<211> 60

<212> DNA

<213> Hepatitis C virus

<220>

<221> misc_feature

<223>/note="antisense"

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<210> 96

<211> 19

<212> DNA

<213> Hepatitis C virus

<400> 96

gtttaaccac tgcattgatg 19

<210> 97

<211> 20

<212> DNA

<213> Hepatitis C virus

<400> 97

gtcccatcga gtgcggctac 20

<210> 98

<211> 45

<212> DNA

<213> Hepatitis C virus

<400> 98

cgtagacatgg tacattccgg acacttggcg cacttcataa gcgga 45

<210> 99

<211> 42

<212> DNA

<213> Hepatitis C virus

<400> 99

tgctcctaac acaatggagc tctggggacga gtcgttcgtg ac 42

<210> 100

<211> 42

<212> DNA

<213> Hepatitis C virus

<400> 100

taccacgacg cgggagctct gttgctcccg aacgcagggc ac

42

<210> 101

<211> 42

<212> DNA

<213> Hepatitis C virus

<400> 101

tgtcgtggtg gggacggagg cctgcctagc tgcgagcgtg gg

42

<210> 102

<211> 48

<212> DNA

<213> Hepatitis C virus

<400> 102

cgttatgtgg cccgggtaga ttgagcactg gcagtcctgc accgtctc

48

<210> 103

<211> 42

<212> DNA

<213> Hepatitis C virus

<400> 103

cagggccggtt ctaggcctcc actgcatcat catatcccaa gc

42

<210> 104

<211> 26

<212> DNA

<213> Hepatitis C virus

<400> 104

ccggaatgta ccatgtcacg aacgac

26

<210> 105

<211> 24

<212> DNA

<213> Hepatitis C virus

<400> 105

gctccattgt gtatgaggca gcgg

24

<210> 106

<211> 23

<212> DNA

<213> Hepatitis C virus

<400> 106

gagctcccgc tgctgggtag cgc

23

<210> 107

<211> 25

<212> DNA

<213> Hepatitis C virus

<400> 107

cctccgtccc caccacgaca atacg

25

<210> 108

<211> 27

<212> DNA

<213> Hepatitis C virus

<400> 108

ctacccgggc cacataacgg gtcaccg

27

<210> 109

<211> 24

<212> DNA

<213> Hepatitis C virus

<400> 109

ggaggcctac aacggccctg gtgg

24

<210> 110

<211> 22

<212> DNA

<213> Hepatitis C virus

<400> 110

ttctatcgat taaatagaat tc

22

<210> 111

<211> 23

<212> DNA

<213> Hepatitis C virus

<400> 111

gccatacgct cacagccgat ccc

23

<210> 112

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 112

Tyr Glu Val Arg Asn Val Ser Gly Ile Tyr His Val Thr Asn Asp Cys
1 5 10 15

Ser Asn Ser Ser
20

<210> 113

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 113

Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr Glu Ala Ala Asp
1 5 10 15

Met Ile Met His Thr
20

<210> 114

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 114

Ala Ala Asp Met Ile Met His Thr Pro Gly Cys Val Pro Cys Val
1 5 10 15

Arg Glu Asn Asn Ser
20

<210> 115

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 115

Pro Cys Val Arg Glu Asn Asn Ser Ser Arg Cys Trp Val Ala Leu
1 5 10 15

Thr Pro Thr Leu Ala
20

<210> 116

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 116

Val Ala Leu Thr Pro Thr Leu Ala Ala Arg Asn Ala Ser Val Pro
1 5 10 15

Thr Thr Thr Ile Arg
20

<210> 117

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 117

Ser Val Pro Thr Thr Thr Ile Arg Arg His Val Asp Leu Leu Val
1 5 10 15

Gly Ala Ala Ala Phe
20

<210> 118

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 118

Leu Leu Val Gly Ala Ala Ala Phe Cys Ser Ala Met Tyr Val Gly
1 5 10 15

Asp Leu Cys Gly Ser
20

<210> 119

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 119

Tyr Val Gly Asp Leu Cys Gly Ser Val Phe Leu Val Ser Gln Leu
1 5 10 15

Phe Thr Ile Ser Pro
20

<210> 120

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 120

Ser Gln Leu Phe Thr Ile Ser Pro Arg Arg His Glu Thr Val Gln
1 5 10 15

Asp Cys Asn Cys Ser
20

<210> 121

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 121

Thr Val Gln Asp Cys Asn Cys Ser Ile Tyr Pro Gly His Ile Thr
1 5 10 15

Gly His Arg Met Ala
20

<210> 122

<211> 20

<212> PRT

<213> Hepatitis C virus

<400> 122

His Ile Thr Gly His Arg Met Ala Trp Asp Met Met Met Asn Trp
1 5 10 15

Ser Pro Thr Thr Ala
20